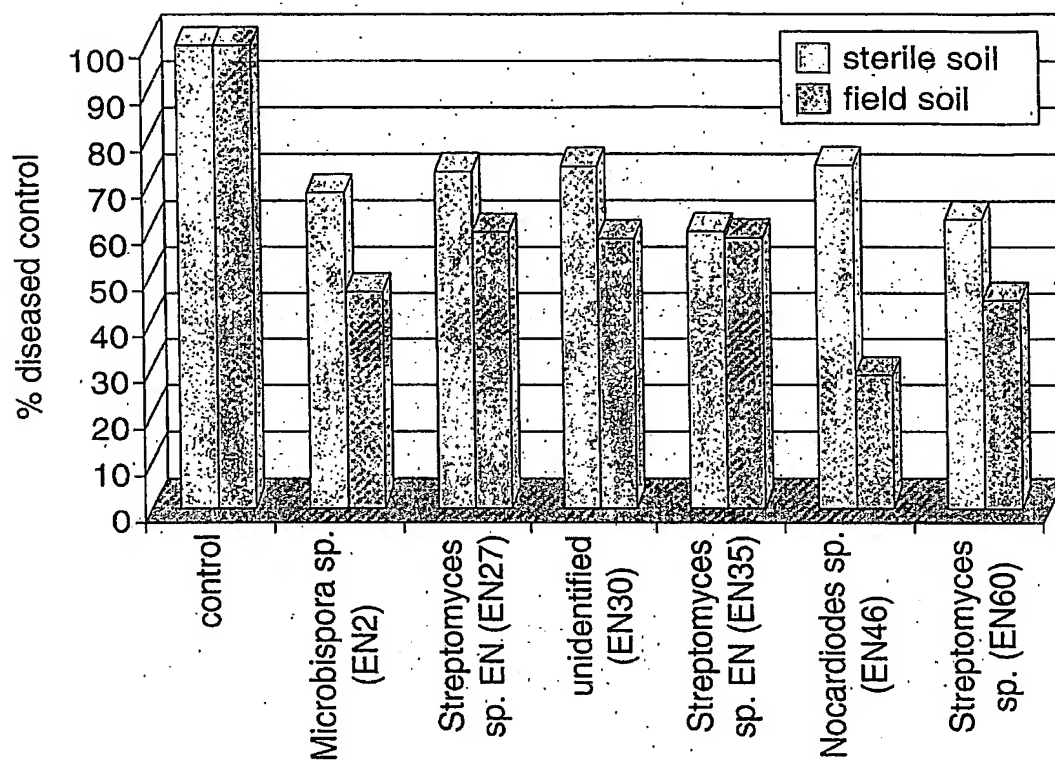


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BEST AVAILABLE COPY

Figure 1

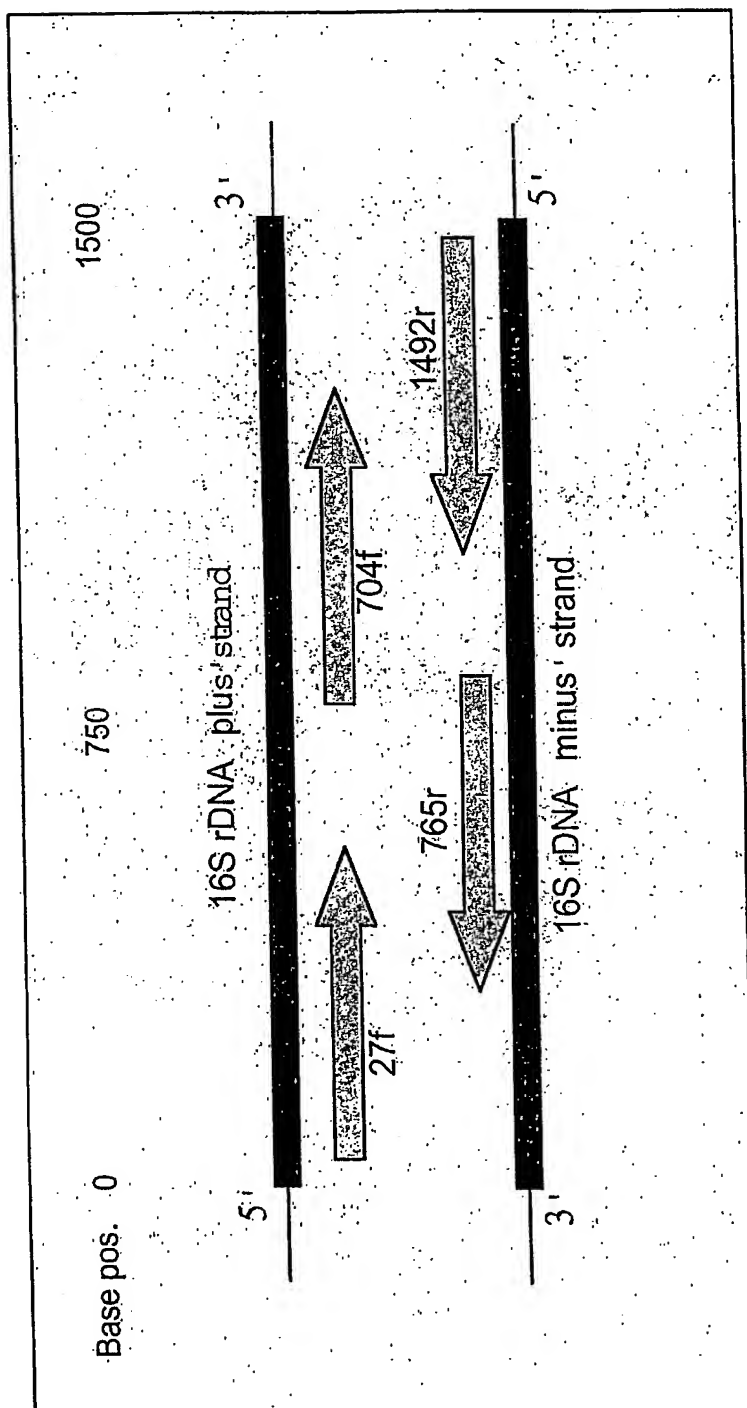
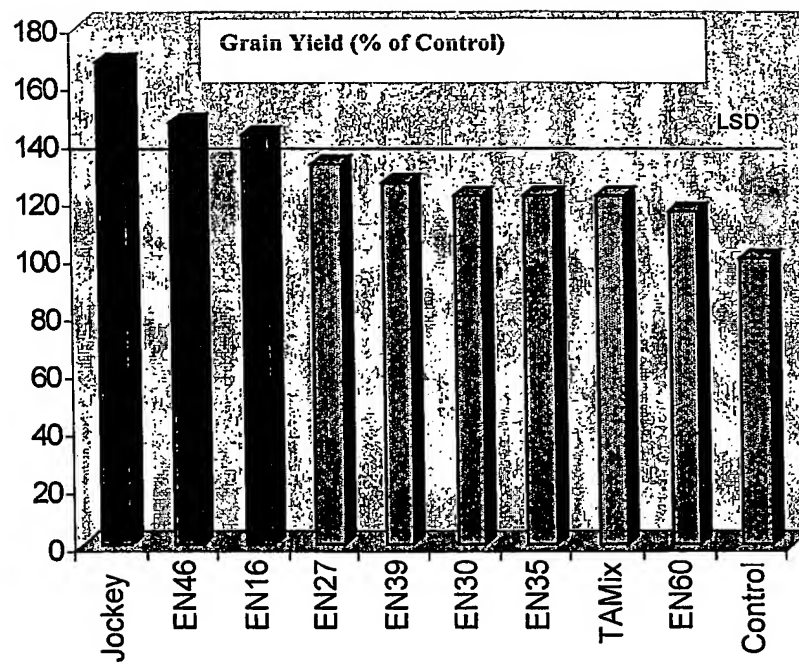
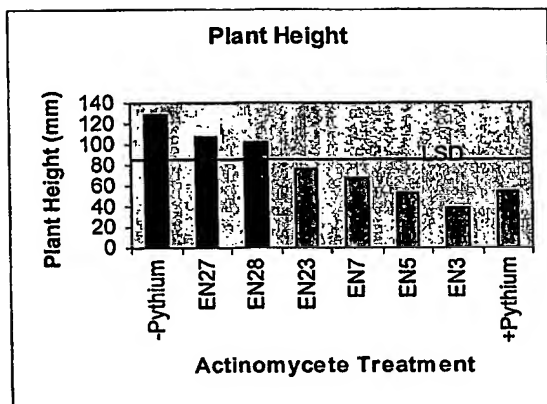
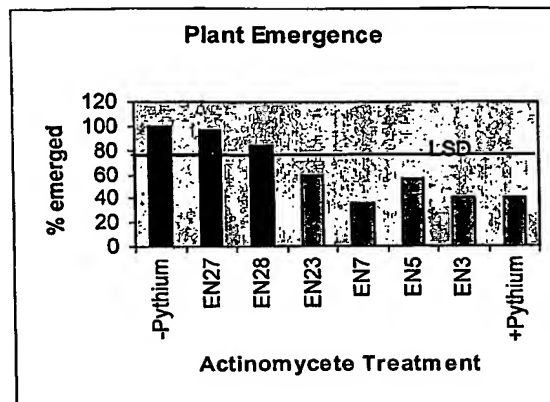


Figure 2

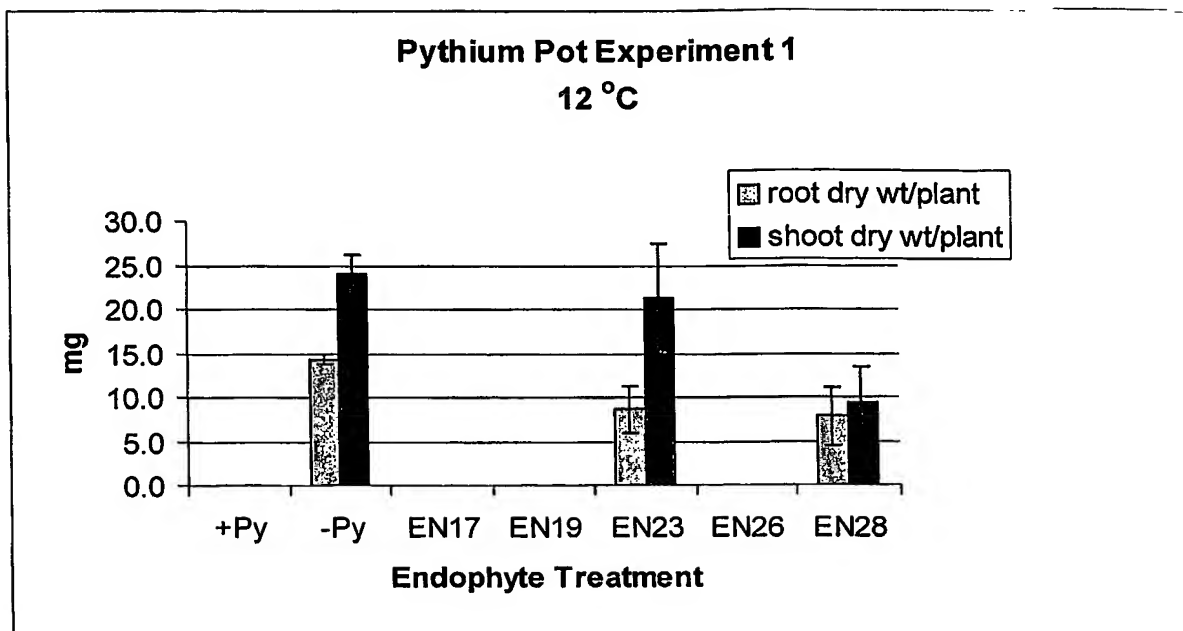
3/30

**Figure 3**

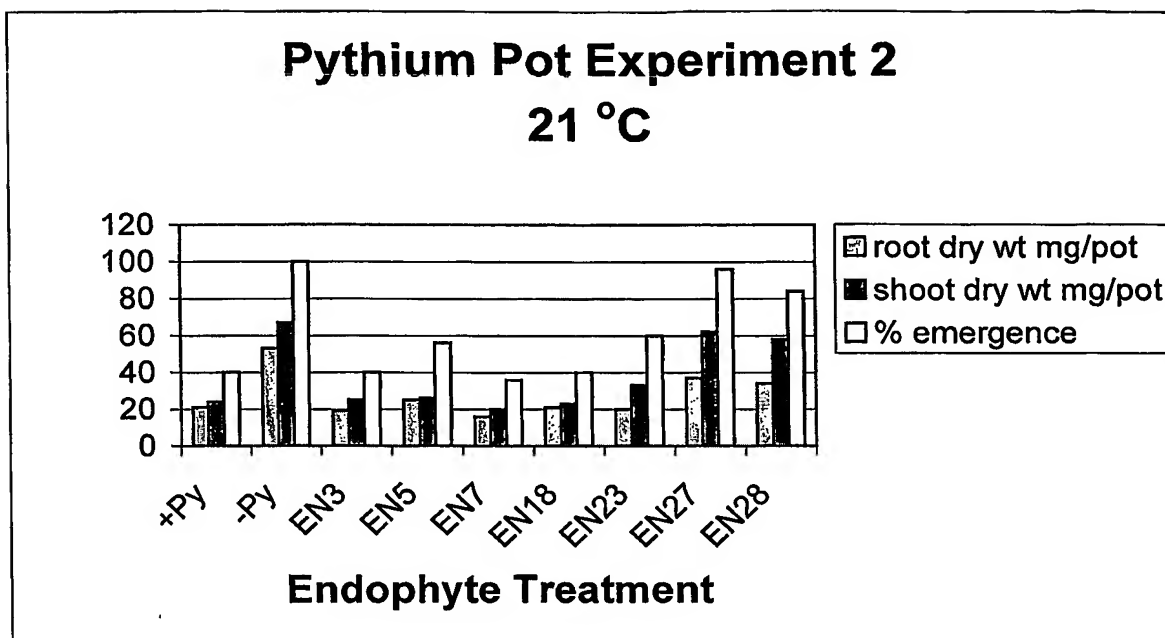
4/30

**Figure 4****Figure 5**

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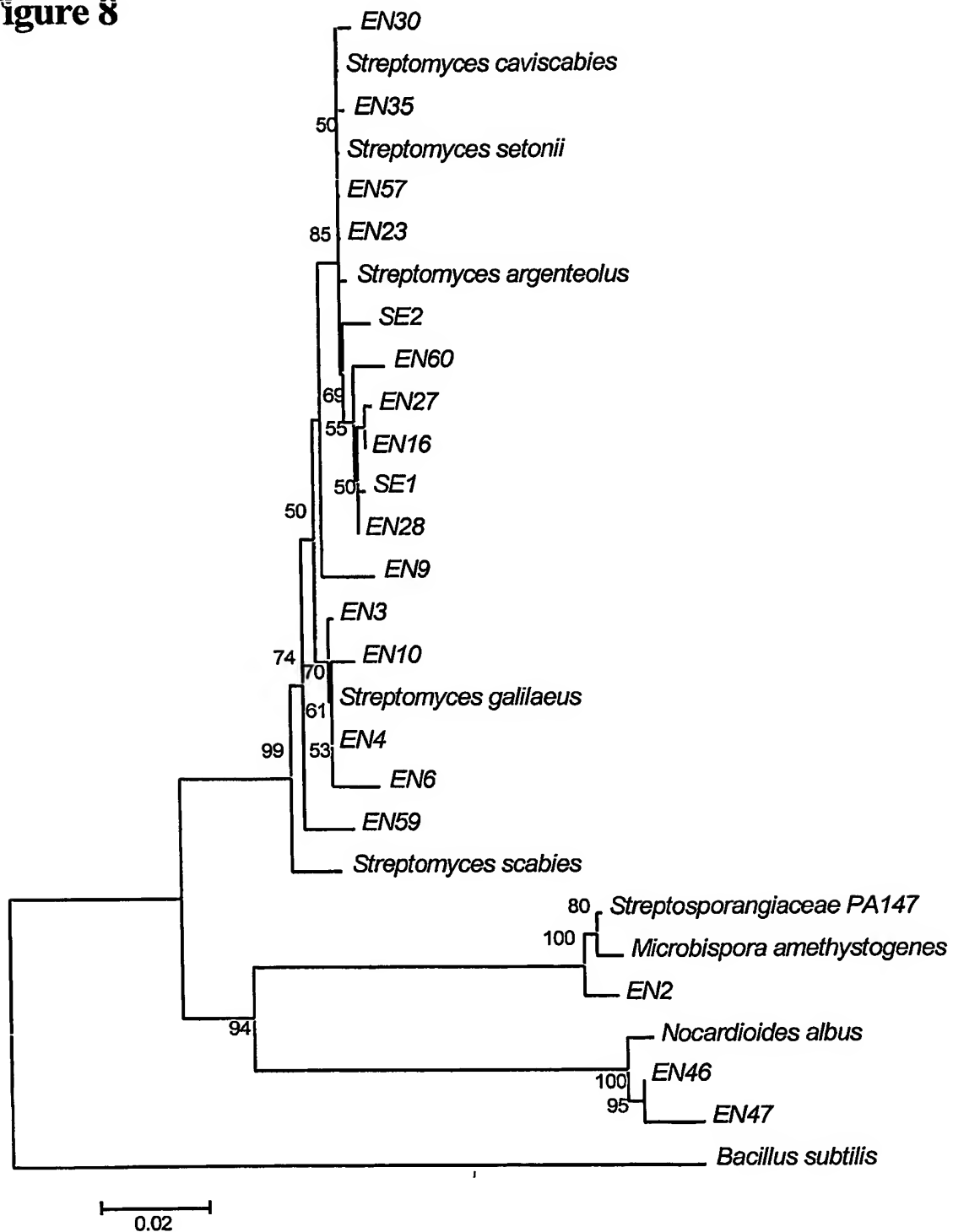
**Figure 6**

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**Figure 7**

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Figure 8



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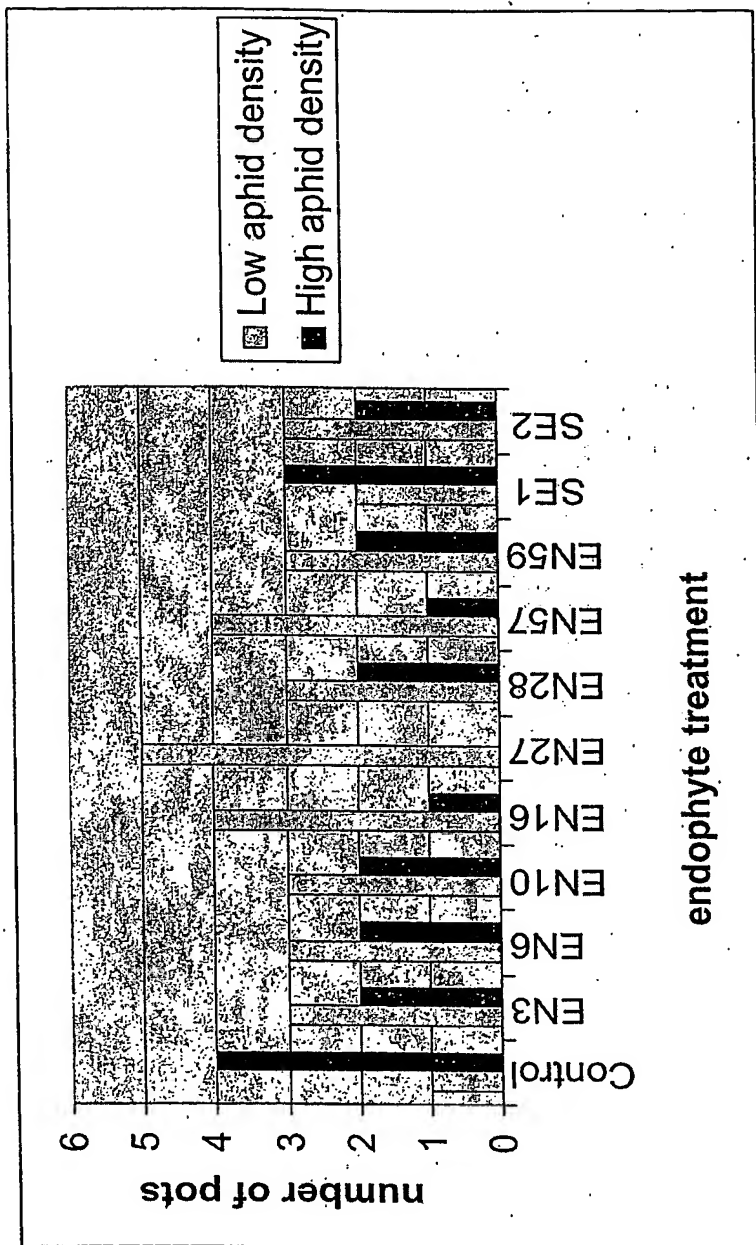


Figure 9

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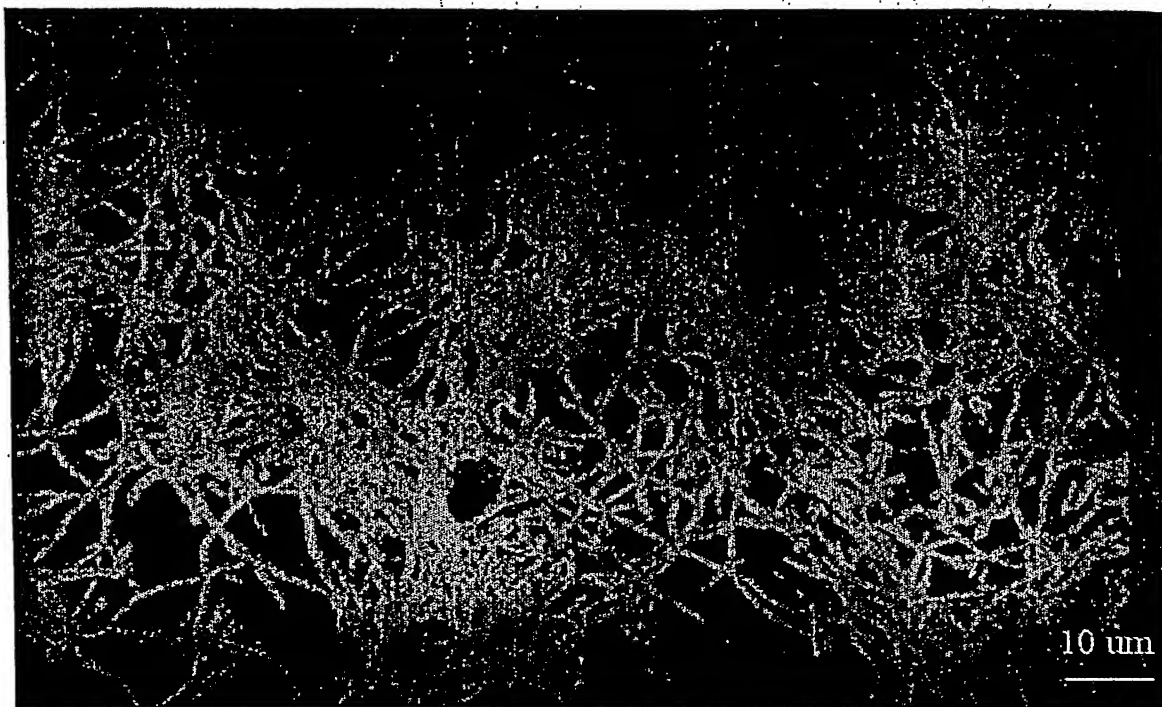
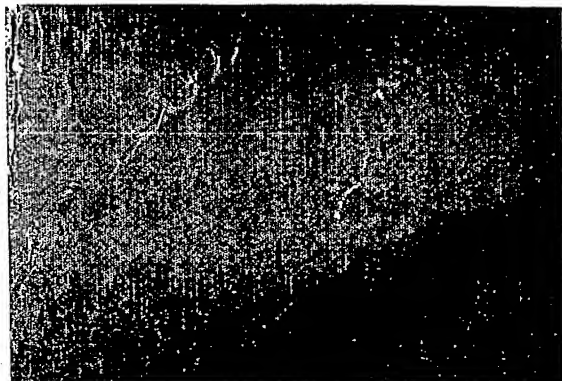


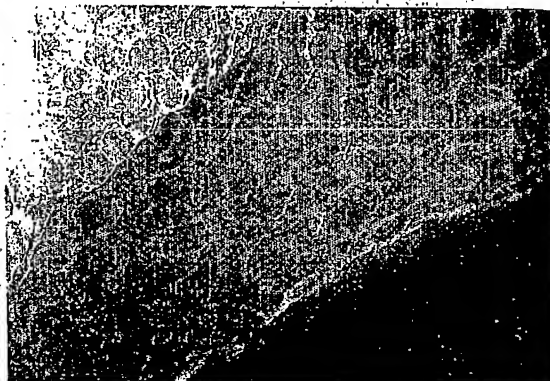
Figure 10

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11.1 Blue excitation/ green emission



11.2 UV excitation/ blue emission



11.3 Image enhanced merge

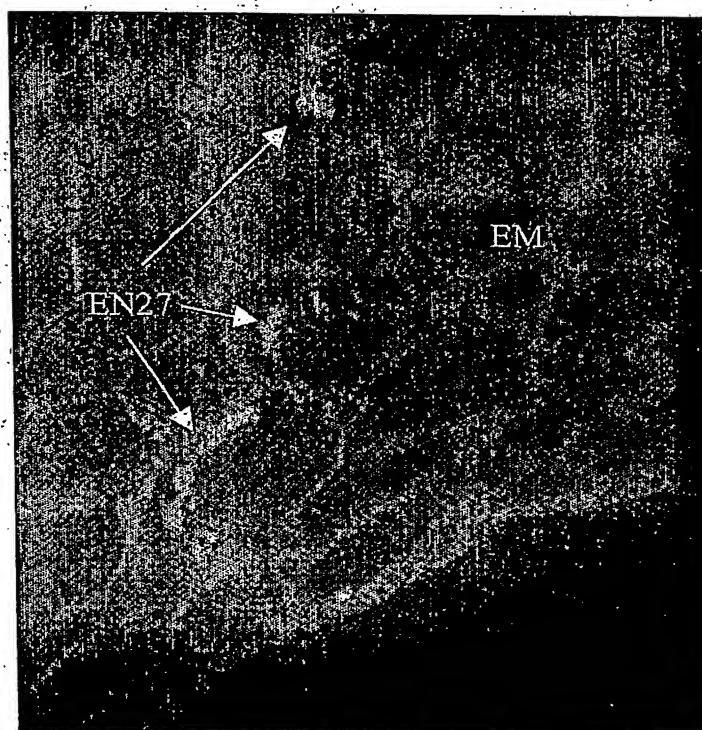
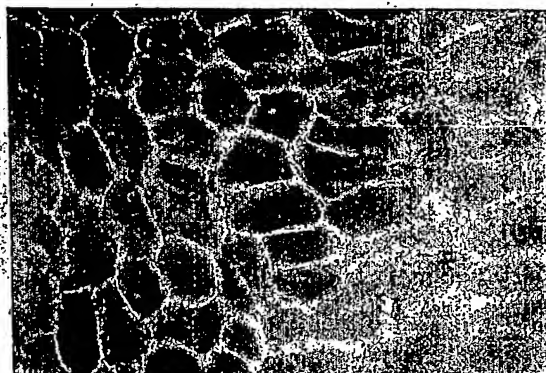
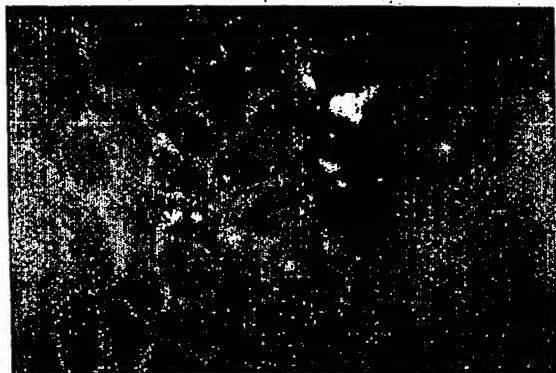


Figure 11

11/30

12.1 Blue excitation/ green emission

12.2 UV excitation/ blue emission



12.3 Image enhanced merge

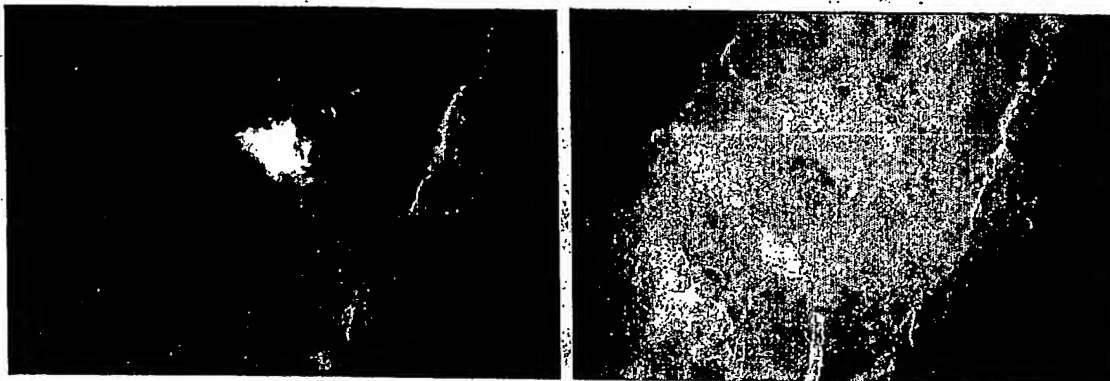


Figure 12

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13.1 Blue excitation/ green emission

13.2 UV excitation/ blue emission



13.3 Image enhanced merge

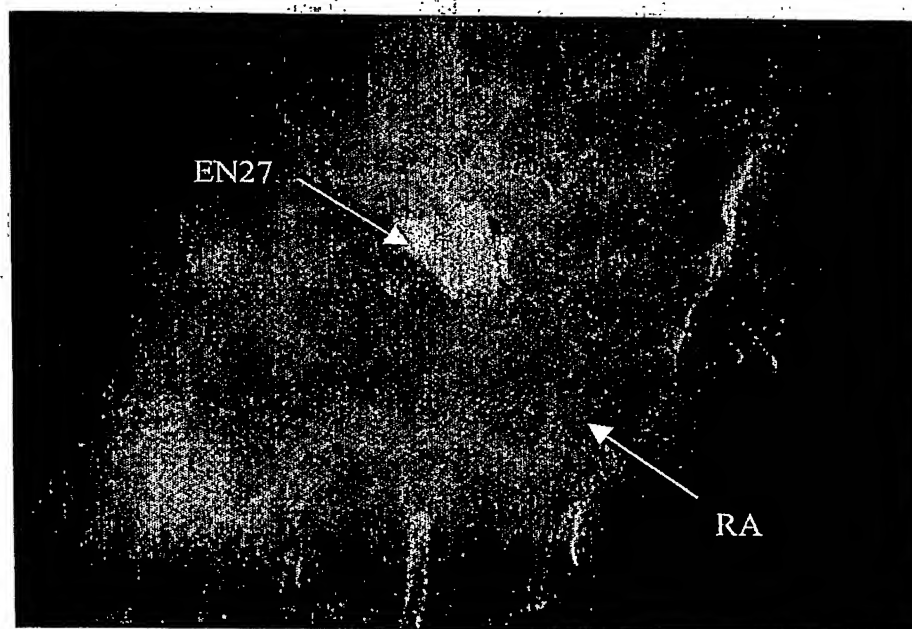
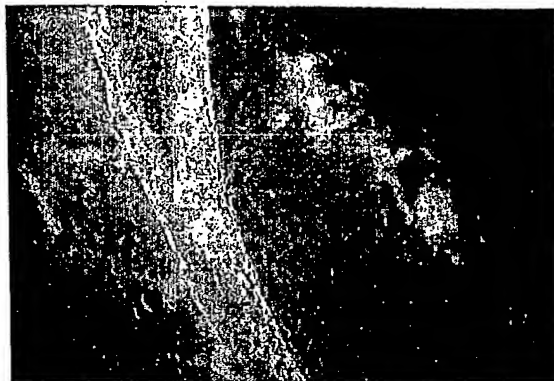


Figure 13

13/30

14.1 Blue excitation/ green emission

14.2 UV excitation/ blue emission



14.3 Image enhanced merge

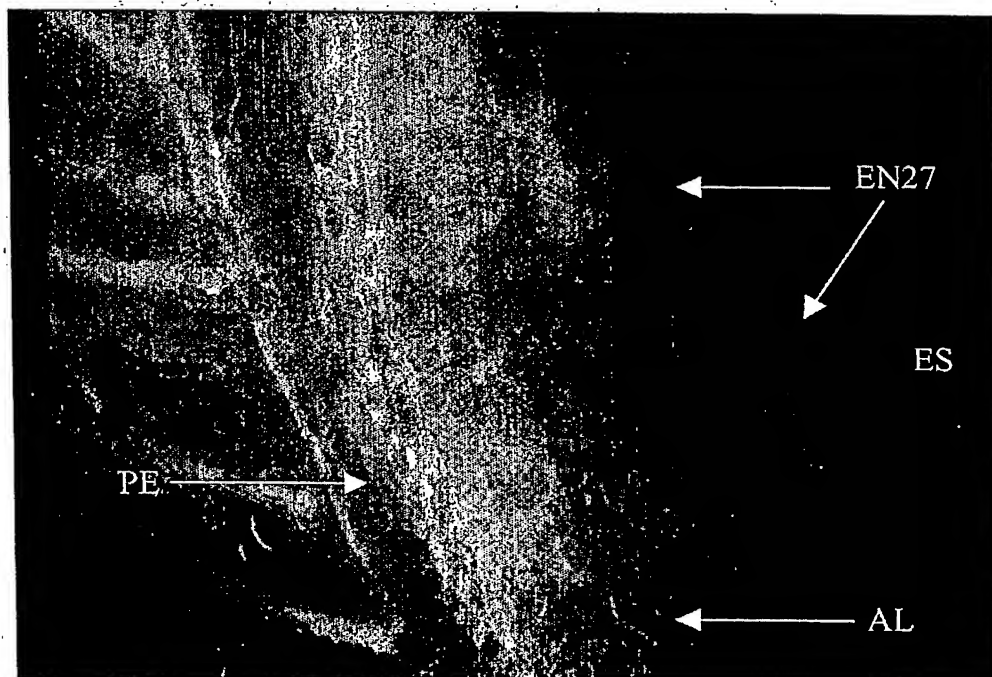


Figure 14

FIGURE 15

EN2

SEQ ID NO:1

CTTAACACATGCAANTCAAGCGGAAAGGCCCTTCGGGGTACTCAANCGGCNAACGGGTGATTAAACACNTGANTAA
CCTGCCCCCTGACTCTGGGATAANCTGGGAAACTGGGTCTAATACCGGATACAACCATTTCTCNCATGGGATGGT
GGTGGAAANTTTTTNCGGTTGGGGATGGGCTCGCGGCCATCACCTTGTGGTGGGGTGATGGCCTACCAAGGCg
ACgAACGGTAgCCCGCTGAgAGGGCgACCGGCCaCaCTGGGACTGAgACaCCGCCgAACTCCTaCgGGAGGCA
gCACTgGGGAATaTTGCCCATGGGCGGAAGCCTGACGCAGNGACGCCGCGTGGGGGATGACGGCCTTNGGGTTGT
AAACCTNTTTTCAGCAGGGACGAAGTTGACGTGTACCTGTAGAGAAGCGCCGGCTAAATANGTGCCAGCAGCCGC
GGTAATANGTAGGGCGCGAGCGTTNTCCGGAATTATTGGGCGTAAAGAGTTTGTAGGTGGCTTGTGCGTTTGCC
GTGAAAGCCCGTGGCTTAANTACGGGTTTGCCTGGATACGGGCAGGCTAGAGGCTGGTAGGGGCAAGCGGAATT
CCTGGTGTAGCGGTGAAATGCGCAGATATCAGGAGGAACACCGGTGGCGAAGGCGGCTTGTGGGCCAgTTcTGA
CGGTGAGGAGCGAAAGCGTGGGGAGCGAACAGGATTAGATACCCTGGTAGTCCACGCTGTAAACGTTGGGCGCTA
GGTGTGGGGGTCTTCCACGATCTCTGTGCCGTAGCTAACGCATTAAGCGCCCCGCTGGGGAGTACGGCCGCAAG
GCTAAACTCAAAGGAATTGACGGGGGCCCGCACAAAGCGGCGGAGCATGTTGCTTAATTCGACGCAACGCGAAGA
ACCTTACCAAGGTTTGACATACACCGGAAACACTCANANATGGGTGCCTCCTTTGGACTGGTGTACAGGTGGTGC
ATGGCTGTNNACACCTCGTGTCTGNAGATGTNGGGTTAAGTCCCGCAACGANCGCAACCCTTGGTTCATGTTG
CCAGCACNCCCTTTGNGGTGGTGGGGACNCATGGGANAATGCCGGGGTCNACTCNGGAGGAAGGTGGGGATGACG
TCAAGTNATCNTGCCCCTTATGTTCTTGNNGTG

EN3

SEQ ID NO:2

GCTGGCGGCGTGCTTAACACATGCAAGTCGAACGATGAACCACTTCGGTGGGGATTAGTGGCGAACGGGTGAGTA
ACACGTGGGCAATCTGCCCTTCACTCTGGGACAAGCCCTGGAAACGGGGTCTAATACCGGATAACACTNCTGCTC
TCATGGGCGAGGGTTAAAAGCTCCGGCGGTGAAGGATGAGCCCGCGGCTATCAGCTTGTGGTGAGGTAATGGC
TCACCAAGGCGACGACGGGTAGCCGGCTGAGAGGGCGACCGGCCACACTGGGACTGAGACACGGCCAGACTCC
TACGGGAGGCGAGCAGTGGGGAATATTGCaCAATGGGCGAAAGCCTGATGCAGCGACGCCGCGTGAGGGATGACG
GCCTTCGGGTGTAAACCTCTTTCAGCAGGGAAGAAGCGAAAGTGACGGTACCTGCAGAAGAAGCGCCGGCTAAC
TACGTGCCAGCAGCCGCGGTAATACGTAgGGCGCAAGCGTTGTCCGGAATTATTGGGCGTAAAGAGCTtGTAgGC
GGCTTGTACGTGCGGTGTGAAAGCCCGGGCTTAACCCcGGGTCTGCATTCGATACGGGCTAgCTAgAGTGTGG
TAGGGGAGATCGGAATTCcTGGTGTAGCGGTGAAATGCgCAGATATCAGGAGGAACACCGGTGGCGAAGGCGGgAT
cTcTGGGCCATTACTGACGCTGAGGAGCGAAAGCGTGGGGAGCGAACAGGATTAGATACCCTGGTAGTCCACGCC
GTAAACGGTGGGAAGTAGGTGTTGGCGACATTCCACGTCGTGCGTGCCGACGCTAACGCATTAAGTTCCCCGCTT
GGGGAGTACGGCCGCAAGGCTAAACTCAAAGGAATTGACGGGGGCCCGCACAAAGCAGCGGAGCATGTGGCTTAA
TTCGACGCAACGCGAAGAACCTTACCAAGGCTTGACATACACCGGAAAGCATCAGAGATGGTGCCCCCTTGTGG
TtCGGTGTACAGGTGGTGCATGGCTGTCTGTCAGCTCGTGTCTGAGATGTTGGGTTAAGTCCCGCAACGAGCGCA
ACCCTTGTCTGTGTTGCCAGCATGCCCTTCGGGGTGATGGGGACTCACAGGAGACCGCGGGGTCAACTCGGAG
GAAGGTGGGGACGACGTCAAGTCATCATGCCCTTATGTCTTGGGCTGCACACGTGCTACAATGGcCGGTACAAA
GAGCTGCGATACCGTGAGGTGGAGCGAATCTCAAAAAGCCGCTCTCAgTTCGGATTGGGGTcTGCAACTCGAcCC
CATGAAGTCGGAGTTGcTAATAATCgCANATCAgCATTGCTGCGGTGAATACGTTcCCGGGCTTGTACAcACCG
CCcGTCACGTcACGAAAGTCGgTAAcACCcGAAgCCGGTGGCCAACCCCTTgTGGGAGGgAGCTGTGCAAGGTGG
GACTGGCGATTG

[FIGURE 15 CONTINUED]**EN16****SEQ ID NO: 7**

GCTTNTTGGTGGGNCNATGGCCTACCAAGGNGAGGACGGNTANCCNGCCTGNGAGGGAGACCGNCCACACTGGGA
ATGNGANACGGCCAGAATCCTACGGGAGGCAGCANNGGGGAANATTGCACAANGGGCGAAAGCCTGATGCAGNG
ANGCCGCGTGAGGGAAGACGGCCTTTGGGTTGTAAACCTNTTTNAGCAGGGAAGAAGCGAAAGTGACGGTACCTG
CAGAAGAAGCGCCGGCTAANTANGTGCCAGCAGCCGCGGTAATANGTAGGGCGCAAGCGTTGTCCGGAATTATTG
GGCGTAAAGAGCTTGTAGGCGGCTTGTGANGTNGGATGTGAAAGCCCGGGGCTTAACCECGGGTTTGCATTTGAT
ACGGGCTAGCTAGAGTGTGGTAGGGGAGATNGGAATTCCTGGTGTAGCGGTGAAATGCGCAGATATCAGGAGGAA
CACCGGTGGCGAAGGCGGATCTCTGGGCCATTACTGACGCTGAGGAGCGAAAGCGTGGGGAGCGAACAGGATTAG
ATACCCTGGTAGTCCACGCCGTAAACGTTGGGAAC TAGGTGTTGGCGACATTCCACGTCGTCGGTGCCGCAGCTA
ACGCATTAAAGTTCCCCGCCTGGGGAGTACGGCCGCAAGGCTAAAACTCAAAGGAATTGACGGGGGCCCCGACAAAG
CAGCGGAGCATGTGGCTTAATTCGACGCAACGCGAAGAACCTTACCAAGGCTTGACATATACCGGAAAGCATCAG
AGATGGTGCCCCCTTGTGGTCGGTATACAGGTGGTGCATGGCTGTCGTGAGCTCGTGTGAGATGTTGGGTT
AAGTCCCGCAACGAGCGCAACCCCTTGTCTGTGTTGCCAGCATGCCCTTCGGGGTGATGGGGACTCACAGGAGAC
TGGGGGTCTGNAACTCGACCCCATGAANTCGGAGTTGCTAATAATCCCAAATTCANCATTGGTGCGGTGAATACT
TCCCGGGCCTGGTACACNACCGCCCGTCAACTCACGAAAGTCGGTNAAACCCGAAACCGGTGGGCCAACCCTTG
TGGGAAGGAAC TGCCNAAGTGGGACTGGCGATTGGGAC

EN23**SEQ ID NO: 10**

ACGAACGCTGGCGCGCTGCTTAACACATGCAAGTCGAACGATGAAGCCGCTTCGGTGGTGGATTAGTGGCGAACG
GGTGAGTAACACGTGGGCAATCTGCCCTTCACTCTGGGACAAGCCCTGGAAACGGGGCTAATAACCGGATAACAC
TCTGTCCCGCATGGGACGGGGTTGAAAGCTCCGGCGGTGAAGGATGAGCCCGCGCCTATCAGCTTGTGGTGGG
GTAATGGCCTACCAAGGCGACGACGGGTAGCCGGCCTGAGAGGGCGACCGGCCACACTGGGACTGAGACACGGCC
CAGACTCCTACGGGAGGCAGCAGTGGGGAATATTGCACAATGGGCGAAAGCCTGATGCAGCGACGCCGCGTGAGG
GATGACGGCCTTCGGGTGTAAACCTCTTTTCAGCAGGGAAGAAGCGAAAGTGACGGTACCTGCAGAAGAAGCGCC
GGCTAACTACGTGCCAGCAGCCGCGGTAATACGTAGGGCGCAAGCGTTGTCCGGAATTATTGGGCGTAAAGAGCT
CGTAGGCGGCTTGTACGTCGGATGTGAAAGCCCGGGGCTTAACCCCGGGTCTGCATTTCGATACGGGCTAgCTAG
AGTGTGGTAGGGGAGATCGGAATTCCTGGTGTAgCGGTGAAATGCGCAGATATCAGGAGGAACACCGGTGGCGAA
GGCGGATCTCTGGGCCATTACTGACgTcTGAGGAGCGAAAGCGTGGGgAgCGAACAGGATTAGATACCCTGgTAG
TCCACGCCGTAAACGTTGGgAACTAGgTGTTGGCGACATTCCACGTCGTGCGTGCCGCAGCTAACGCATTAAAGTT
CCCCGCTGGGGAGTACGGCCGCAAGGCTAAAACTCAAAGGAATTGACGGGGGCCCCGACAAAGCAGCGGAGCATG
TGGCTTAATTGACGCAACGCGAAGAACCTTACCAAGGCTTGACATATACCGGAAAGCATCAGAGATGGTGCCCC
CCTTGTGGTCGGTATACAGGTGGTGCATGGCTGTCGTGAGCTCGTGTGAGATGTTGGGTAAAGTCCCGCAAC
GAGCGCAACCCCTTGTCTGTGTTGCCAGCATGCCCTTCGGGGTGATGGGGACTCACAGGAGACTGCCGGGGTCAA
CTCGGAGGAAGGTGGGGACGACGTCAAGTCATCATGCCCTTATGTCTTGGGCTGCACACGTGCTACAATGGCCG
GTACAATGAGCTGCGATGCCGCGAGGCGGAGCGAATCTCAAAAAGCCGGTCTCAGTTCGGATTGGGGTCTGCAAC
TCGACCCCATGAAGTCGGAGTTGCTAGTAATCGCAGATCAGCATTGCTGCGGTGAATACGTTCCCGGGCCTTGTA
CACACCGCCCGTCACGTCACGAAAGTCGGTAACACCCGAAGCCGGTGGCCCAACCCTTGTGGGAGGGAGCTGTGCG
AAGGTGGGACTGGCGATTG

[FIGURE 15 CONTINUED]

EN27

SEQ ID NO:12

TTAANACATGCAANTCGAACGATGAACCCNGTTTCGGTGGTGGATTAGTGGCGAACGGTGAGTAANANGTGGGCA
ATTTGCCCTTCATTTTGGACAAGCCCTGGAAACGGGTTTAATACCGGATAACATTTTNTCCCGCATGGGANGGGG
TTGAAAGNTCCGGCGGTGAAGGATGAGCCCGCGGCCATNAGCTTGTTGGTGGGGTAATGGCCTACCCAAGGGAG
ACGGGTAGCCGGCCTGAGAGGGCGACCGGCCACACTGGGAATGAGANACGGCCCAGAATCCTACGGGAGGCAGCA
GTGGGGAATATTGCACAATGGGCGAAAGCCTGATGCAGCGANGCCGCGTGAGGGATGACGGCCTTNGGGTTGTAA
ACCTTTTTNAGCAGGGAAGAAGCGAAAGTGACGGTACCTGCAGAAGAAGCGCCGGCTAAATAAGTGCCAGCAGCC
GCGGTAATAAGTAGGGCGCAAGCGTTGTCCGGAATTATTGGGCGTAAAGAGCTTGTAGGGCGCTTGTCANGTNGG
ATGTGAAAGCCCGGGGNTTAACCCCGGGTTTGCATTTGATACGGGCTAGNTAGAGTGTGGTAGGGGAGATNGGAA
TTCCTGGTGTAGCGGTGAAATGCGCAGATATCAGGAGGAACACCGGTGGCGAAGGCGGATCTCTGGGCCATTACT
GACGCTGAGGAGCGAAAGCGTGGGGAGCGAACAGGATTAGATACCCTGGTAGTCCACGCCGTAAACGTTGGGAAC
TAGGTGTTGGCGACATTCCACGTCGTCGGTGCCGACGCTAACGCATTAAGTTCCCCGCTGGGGAGTACGGCCGC
AAGGCTAANACTCAAAGGAATTGACGGGGGCGCCNACAAGCAGCGGANCATGTGGCTTAATTTCGACGCANCGCGA
AGAACCTTACCAAGGCTTGACATATACCGGAAAGCATCAGAGATGGTGCCCCCTTGTTGGTCGNTATACANGTGG
TGCATGNCTGTCTCACCTCGTGTCTGTGAGATGTTGGGTAAAGTCCCGCAACGAGCGCNACCCTTGNTCTGTGTT
GNCANCATGCCCTTCGGGGNTGATGGGGACTCACAGGANACTGNCCGGGGTCAACTCCGGANGAAGGTGGGTGAC
GAAGTCAAGGTTCATCATGNCCCCTTATGTCTTGGTGTCTGCACACGTGC

EN28

SEQ ID NO:13

TTCGGNGGTGGANTAGNGGCGNACGGGNGACCAACANGNGGGCAATCCCCCCTTCANTTTNGGACAACCCCTGGA
AACGGGTTNTAATACCGGATAACANTTTNTCCCGCATGGGANGGGGTGAAAGCTCCGGCGGTGAAGGATGAGC
CCGCGGCCATCAGCTTGTTGGTGGGGTAATGGCCTACCAAGGCGACGACGGGTAGCCGGCCTGAGAGGGCGACC
GGCCACACTGGGANTGAGANACGGCCCAGAATCCTACGGGAGGCAGCAGTGGGGAATATTGCACAATGGGCGAAA
GCCTGATGCAGCGACGCCGCGTGAGGGATGACGGCCTTCGGGTGTAAACCTTTTTTCAGCAGGGAAGAAGCGAAA
GTGACGGTACCTGCAGAAGAAGCGCCGGCTAAATANGTGCCAGCAGCCGCGGTAATANGTAGGGCGCAAGCGTTG
TCCGGAATTATTGGGCGTAAAGAGNTTGTAGGCGGCTTGTCANGTCGGATGTGAAAGCCCGGGGCTTAACCCCGG
GTTTGCATTTCGATACGGGCTAGCTAGAGTGTGGTAGGGGAGATCGGAATTCCCTGGTGTAGCGGTGAAATGCGCAG
ATATCAGGAGGAACACCGGTGGCGAAGGCGGATcTcTGGGCCATTACTGACGcTGAGGAGCGAAAGCGTGGGgAG
CGAACAGgAATTAGATACCCTGgTAGTCCACGCCGTAAACGTTGGgAAcTAGgTGTGGcGACATTCCACGTcGT
CGgTGCCGCGAGCTAACGCATTAAGTTCCCCGCTGGGGAGTACGgCCCGCAAGGCTAAAACCTCAAAGGAATTGAC
GGGGgCCCGCACAAAGCAGCGGAGCATGTGGCTTAATTTCGACGCAACGCGAAGAACCTTACCAAGGCTTGACATAT

ACCGGAAAGCATCAGAGATGGTGCCCCCTTGTTGGTTCGGTATACAGGTGGTGCATGGCTGTCTGCTCAGCTCGTGTC
GTGAGATGTTGGGTTAAGTCCCGCAACGAGCGCAACCCCTTGgTTCTGTGTGgCCAGCATGCCCTTCGGGGTGAT
GGGGACTCACAGGAGACTGgCCGGGGTCAACTCGGAGGAAGGTGGGGACGACGTCAAGTCATCATGCCCTTATG
TCTTGGGgCTGCACACGTGCTACAATGGCCGGTACAATGAGCTGCGATGCCGCGAaGGCGGAGCGAATCTCAAAA
AaGCCGGTCTCAGTTTCGGATTGGGGTCTGCAACTCGACCCCATGAAGTCGGAGTTGCTAGTAATCGCAGATCAGC
ATTGCTGCGGTGAATACGTTCCCGGGCCTTGACACACCGCCCGTCACGTACGAAAGTCGGTAACACCCGAAGC
CGGTGGTCCAACCCCTTGTTGGGAGGGAGCTGTCGAAGGTGGGACTGGCGATTGG

[FIGURE 15 CONTINUED]

EN46

SEQ ID NO:16

ATGCAAGTCGAGCGGAAAGGCCCTTCGGGGTACTCGAGCGGCGAACGGGTGAGTAACACGTGAGTTAATCTGCCC
CAGGCTCTGGATACCCACCGGAAACGGTGATTAATACCGAATACGACAACCGATTTGCATGATCTGGTgGTGNA
AAGTTTTTTCGGCTGGGATGTGCTTCGCGGCCCTATCAGCTTGTGGTGAGGTAATGGCTCACCCAAGGCTTCGAC
GGTAGCCGGCCTGAGAGGGTGACCGNCCACACTGGGACTGAGACACGGCCAGACTCCTACGGGAGGCAGCAGTG
GGGAATATTGGACAATGGGCGGAAGCCTGATCCAGCAACGCCGCGTGAGGGATGACGGCCTTCGGGTGTAAACC
TCTTTTCAGCACAGACGAAGCGCAAGTGACGGTATGTGCAGAAGAAGGACCGGCCAACTACGTGCCAGCAGCCGCG
GTAATACGTAGGGTCCGAGCGTTGTCCGGAATTATTGGGCGTAAAGGGCTCGTAGGCGGTCTGTGCGCTCGGGAG
TGAAAACAGGTGCTTAACACCTGGCCTGCTTTTCGATACGGGCAGNCTAGAGGTACNCAGGGGAGAATGGAATTC
CTGGTGTAGCGGTGAAATGCGCAGATATCAGGAGGAAACACCGGTGGCGAAGNCGGTTCTCTGGGAGTATCCTGA
CGCTGAGGAGCGAAAGTGTGGGGAGCGAACAGGATTAGATACCCCTGGTAGTCCACACCGTAAACGTTGGGCGCTA
GGTGTGGGACACATTCCACGTGTTCCGTGCCGCGAGCTAACGCATTAANCGCCCCGCTGGGGAGTACGGCCGCAA
NGCTAAAACTCANAGGAATTGACGGGGGGCCGACAAAGCGGCGGAgCATGCGGATTAATTCGATGCAACGCGAAG
AACCTTACCTGGGTTTGACATACACCGGAAAGCCGTACAGATACGGCCCCCTTTAGTCgGTGTaCAGGTGGTGCA
TGGCTGTCTgCgCtCgCTGTCTGTGAGATGTtCGGGTTAAGTCCCGCAACGAGCGCAaCCCTCgTCCTATGTTGC
CaGCAATTTCGGTTGgGGACTCATAGGAgACTGCCgGGGTCaACTCGGAGGAAGGTGGGGATGACGTCAAGTCATC
ATGCCCCCTATGTCCAGGGCTTCACGCATGCTACAATGGCCGGTACAAAGGGCTGCGATCCCGTGAGGGTGAGCG
AATCCCAAAAAGCCGGTCTCAGTTCGGATTGGGGTCTGCAACTCGACCCCATGAAGTCGGAGTCGCTAGTAATCG
CAGATCAGCAACGCTGCGGTGAATACGTTCCCGGGCCTTGTAACACCGCCCCGTACGTACAGAAAGTCGGCAAC
ACCCGAAGCCANTGGCCCACTCGTAAGAGAGGGAGCTGT

EN60

SEQ ID NO:18

ATGCAAGTNGAACGATGAANCCNTTTGGGGTGGATTAGTGGCGAACGGGTGAGTAANANGTGGGCAATTTGCCCT
TCAATTTGGGACAAGCCCTGGAAACGGGGTNTAATACCGGATAACANTNTGTCCCGCATGGGACGGGGTTAAAAG
CTCCGGCGGTGAAGGATGAGCCCGCGGCCTATNAGCTTGTGGTGGGGTGATGGCCTACCAAGGCGACGACGGGT
AGCCGGCCTGAGAGGGCGACCGGCCACACTGGGACTGAGACACGGCCCAGACTCCTACGGGAGGCAGCAGTGGGG
AATATTGCACAATGGGCGAAAGCCTGATGCAGCGACGCCGCGTGAGGGATGACGGCCTTCGGGTGTAAACCTTT
TTCAGCAGGGGAAGAAGCGAAAGTGACGGTACCTGCAGAAGAAGCGCCGGCTAAATANGTGCCAGCAGCCGCGTA
ATANGTAGGGCGCAAGCGTTGTCCGGAATTATTGGGCGTAAAGAGTTTGTAGGCGGCTTGTACGTNGGATGTGA
AAGCCCGGGGCTTAACCCGGGTTTGCATTGATACGGGCTAGCTAGAGTGTGGTAGGGGAGATCGGAATTCCTG
GTGTAGCGGTGAAATGCGCAGATATCAGGAGGAACACCGGTGGCGAAGGCGGaTcTcTGGGCCATTACTGACGNT
GAGGAGCGAAAGCGTGGGGAGCNAACAGNATTAGATACCCCTGGTAGTCCAAGCCGTAAACGTTGGGAACANGTG
TTGGCGACATTCACGTGTCNNTGCCGCANCTAACGCATTAAGTTCCCCGCTGGGGAGTACGGCCGCAAGGCT
AANACTCAAAGGAATTGANGNNGGCCCCGACAAAGCAGCGGAGCATGTGGCTTANTTCNACGCANCGCGAAGAACC
TTACCAAGGTTTGGCATATAcCGGAAAgCaTCAGAgATgGTGCCCCCTtGTGGTCGGTATACAGgTGGTGCNTG
GCTGTCTGTCAGCTCGTGTCTGACAtGtTGGTTAAgTCCCGTCAaCGAGgCGCAACCCTTGTTNTGTGTNGCCAG
CATGCCCTTCGGGGTGATGGGGACTCACAGGAGACTGCCGGGGTCAACTCGGAGGAAGGTGGGGACGACGTCAAG
TCATCATGCCCTTATGTCTTGGGCTGCACACGTGCTACAATGGCCGGTACAATGAGCTGCGATGCCCGGAGGCG
GAGCGAATCTCAAAAAGCCGGTNTCAGTTCGGATTGGGGTCTGCAACTCGACCCCATGAAGTCGGAGTTGCTAGT
AATCGCAGATCAGCATTGCTGCGGTGAATACGTTCCCGGGCCTTGTAACACCGCCCCGTACGTACAGAAAGTCG
GTAACACCCGAAGCCGNTGG

[FIGURE 15 CONTINUED]

PM87

SEQ ID NO:24

GGCCCAGANATCCGNCTTCGCCACCGGTGTTCCCTCCTGAATATCTGCGCATTTACCGCTACACCAGGAATTCCG
ATCTCCCCTACCACACTCTAACTAGCCCGTATCGAATGCAGACCCGGGGTTAAGCCCCGGGCTTTCACATCCGAC
GTGACAAGCCGCCTACGAGCTCTTTACGCCCAATAATTCGGGACAACGCTTGCGCCCTACGTATTACCGCGGCTG
CTGGCACGTAATTAGCCGGCGCTTCTTCTGCAGGTACCGTCACTTTTCGCTTCTTCCCTGCTGAAAGAGGTTTACA
ACCCGAAGGCCGTCATCCCTCACGCGGCGTCGCTGCATCAGGCTTTGCCCCATTGTGCAATATTCCCCACTGCTG
NCTCCCGTANGAGTCTGGGCCGTGTCTCAGTCCCAGTGTGGCCGGTCGNCCTCTCAGGCCGGCTACCGTCGTCGC
CTTGGTAGGCCATTACCCACCAACAAGCTGATANGCCGNGGGCTCATCCTTCANCGTCGGAGCTTCAANCCCG
TCCATGCGGGACAGAGTGTATCCGGTATTANACCCCGTNTCAGGGCTTGTCANAGTGAAGGGCAGATNGCCAC
GTGTTATCACCGTTCGCCACTAATNACANCGAACGGCTTATCGTNCGACTGCATGTGTTAACACNCGCAGCGTT
CGTCTGAGCCAG

FIGURE 16

EN5

SEQ ID NO:3

GTAATGGCCCANAAAAACCGCCTTCGCCACCGGTGTTCTCCTGATATCTGCGCATTTACCGCTACACCAGGAAT
TCCNATCTCCCCTACCACACTCTAGCTAGCCCGTATCNAATGCAAACCTCGGGGTTAAGCCCNAGCTTTCACATC
CGACGTGACAAGCCGCCTACAANCTCTTTACGCCCAATAATTCCGGANAACGCTCGCACCCCTACNTNTTACCGCG
GCTGCTGGCNCGTNTTTAGCCGGTGCTTCTTCTGCAGGTACCGTCACTTTCGCTTCTTCCCTGCTNAAAAAGGTT
TACAACCCCTANGGCCGT

EN6

SEQ ID NO:4

TGAGGGATGACGGCNTTCGGGGTTGTAAACNTTNTCACCAGGGAAGAAGCGAAAGTGNCGGTACCTGCAGAAGA
AGCGCCGNCTAACTACGGGCCAGCATCCGCGGTAATACGTAGGGCGCAATCGTTGTCCGGAATTANTGGGCGTAA
AGAGNTCGTAGGCGGCTTATCACGTCGGGTGTGAAAGCCCGGGGCTTAAGCCCCGGGTCTGCATTCGATACGGGC
TAGCTAGANTNTGNTAGGGGAGATCGGAATTCCTGGTGTAGCGGTGAAATGCGCAGATATCAGgAGGAACACCGg
TGGCGAAGGCGGATCTcTGGGCCATTACTGACGCTGAGGAGCGAAAGCGTGGGGAGCGAACAGGATTAGATACCC
TGGTAGTCCACGCCGTAAACGGTGGGAACCTAGGTGTTGGCGACATTCCACGTCGTCGGTGCCGACGCTAACGCAT
TAAGTTCCTCCGCTGGGGAGTACGGCCGCAAGGCTAAAACTCAAAGGAATTGACGGGGGCCCGCACAAAGCAGCGG
AGCATGTGGCTTAATTCGACGCAACGCGAAGAACCTTACCAAGGCTTGACATACACCGGAAAGCATCAGAGATGG
TGCCCCCTTGTGGTGGTGTACAGGTGGTGCATGGCTGTCTGTCAGCTCGTGTCTGAGATGTTGGGTAAAGTCC
CGCAACGAGCGCAACCCCTTGGTCTGTGTTGCCAGCATGCCCTTCGGGGTGATGGGGACTCACAGGAGAACGCCG
GGGTCAACTCGGAGGAAGGTGGGGACGACGTCAAGTCATCATGCCCTTATGTCTTGGGCTGCACACGTGCTACA
ATGGCAGGTAAATGAGCTGCGATACCGTGAGGTGGAGCGAATCTCAAAAAGCCTGTCTCANTTCGGATTGGGGT
CTGNAANTCGACCCCATGAAAGTCGGAGTTGCTAATTATCCCAGATCAACATTGCTGGCGGTGAATACGTTCCCG
GGGCCCTTGGTAAACACCGCCCGTCAANGTNAAGAAAGTCGGGTAACACCCGAAANCCGGTGGGCCAANCCCT

EN7

SEQ ID NO:5

CCGCCTTCGCCACCGGGTGTTCCTCCTGATATCTGCGCATTTACCGCTACACCAGGAAATTCNATCTCCCCTA
CCACACTCTANCTANCCCGTATCGAATGCAAAACCCGGGGTTAANCCCCGGGCTTTCACACCCGACNTGACAAGCC
GCCTACAAACTCTTTACGCCCAATAATTCCGGACAACGCTTGCGCCCTACNTATTACCGCGGTGCTGGCACNTA
TTAGCCGCGCTTCTTCTGCAGGTACCGTCACTTTCGCTTCTTCCCTGCTGAAAAAGGTTTACAACCCGAAGGC
CGTCATCCCTCAGCGGCGTGCCTGCATCAGGCTTTCGCCCATTGTGCAATATTCCCACCTGCTGCCTCCNTAG
GAATCTGGGCGGTGTCTCAATCCAGTGTGGCCGCTCCCTCTCNGCCGGCTACCGTCNTCCCTTGGTNACCATT
ANCTCACCAACAACCTGATAGNCGCGGGCTCATCTTCACGCGGGAACCTTCAACCACC

EN9

SEQ ID NO:6

GGCGGCGTGCTTAACACATGCAAGTCGAACGATGAAGCCCTTCGGGGTGGATTAGTGGCGAACGGGTGAGTAACA
CGTGGGCAATCTGCCCTTCACTCTGGGACAAGCCCTGGAAACGGGGTCTAATACCGGATACGATTCTGgGAGGCAT
CTCCTGGTaCTGGAAAGCTCCGGCGGTGAAGGATGAGCCCGCGGCcTATCAGCTTGTTGTGGGTAATGGCCTACC
AAGGCGACGACGGgTAGCCGGCCTGAGAGGGCGACCGGCCACACTGGGACTGAGACACGGCCCAGACTCCTACGG
GAGGCAGCAGTGGGGAATATTGCACAATGGGCGAAAGCCTGATGCAGCGACGCCGCTGAGGGATGACGGCCTTC
GGGTTGTAAACCTCTTTCAGCAGGGAAGAAGCGAGAGTGACGGTACCTGCAGAAGAAGCGCCGGCTAACTACGTG
CCAGCAGCCGCGGTAATACGTAGGGCGCAAGCGTGTCCGGAATTATTGGGCGTAAAGAGCTCGTAGGCGGCTTG
TCACGTCCGGGTGTGAAAGCCCGGGGCTTAACCCCGGGTCTGCATCCGATACGGGCAGGCTAGAGTGTGGTAGGGG
AGATCGGAATTCTTGGTGTAGCGGTGAAATGCGCAGATATCAGGAGGAACACCGGTGGCGAAGGCGGATCTCTGG
GCCATTACTGACGCTGAGGAgCgAAAGCGTGGGGAGCCaACAGGATTAGATACCCTGGTAGTCCACGCCGTAAC
GTTGGAAGTGGTGTGGCGACATTCCACGTCGTCGGTGCCGACGCTAACGCATTAAGTTCCTCCGcTGGGGAGT
ACGGCCGCAAGGCTAAAACTCAAAGGAATTGACGGGGGCCGACAAGCAGCGGAGCATGTGGCTTAATTTCGACG
CAACGCGAAGAACCTTACCAAGGCTTGACATATACCGGAAAGCGCCAGAGATGGTGCCCCCTTGGTGGTAT
ACAGGTGGTGCATGGCTGTCTGTCAGCTCGTGTCTGAGATGTTGGGTAAAGTCCCGCAACGAGCGCAACCCCTGT

[FIGURE 16 CONTINUED]

CCTGTGTTGCCAGCATGCCCTTCGGGGTGATGGGGACTCACAGGAGACCGCCGGGTCAACTCGGAGGAAGGTGG
GGACGACGTCAAGTCATCATGCCCCTTATGTCTTGGGCTGCACACGTGCTACAATGGCCGGTACAAAGAGCTGCG
ATGCCGTGAGGCGGAGCGAATCTCAAAAAGCCGGTCTCAGTTCGGATTGGGGTCTGCAACTCGACCCCATGAAGT
CGGAGTTGCTAGTAATCGCAGATCAGCATTGCTGCGGTGAATACGTTCCCGGGCCTTGACACACCGCCCGTCA
GTCACGAAAGTCGGTAACACCCGAAGCCGGTGGCCCAACCCCTCGGGGAGGAGCTGTCTGAAGGTGGGAC

EN17**SEQ ID NO:8**

CCGCCTTCGCCACCGGTGTTCTCCTGATATCTGCGCATTTACCGCTACACCAGGAATTCC
NATCTCCCCTACCACACTCTAGCTAGCCCGTATCAAATGCAAACCCGGGGTTAAGCCCCGGGCTTTC
ACATCCNACGTGACAAGCCGCTACAANCTCTTTACGCCCAATAATTCCGGACAACGCTTTCGCCCT
ACNTATTACCGCGGCTGCTGGCACNTATTTAGCCGGCGCTTCTTCTGCAGGTACCGTCACTTTTCGCT
NCTTCCCTGCTGAAANAGGTTTACAACCCAAAGGCCNTCATCCCTCNCGGGNTCNTTGCNTCNGGC
TTNCNCCCATTGTTCAANNNTCCCACTGCTNCCTCCCCTCGGAATCTGGGCCGNTGTCTCATTCCCN
TTNTGGCCGGTCCCCCTCNCAGGCCNGCTACCC

EN19**SEQ ID NO:9**

CTCAGCGTCNGTAATGGCCCAAAAACCGCCTTCGCCACCGGTGTTCTCCTGATATCTGCGCATTTACCGCTAC
ACCAGGAATTCCNATCTCCCCTACCACACTCTAGCTAGCCCGTATCNAATGCAAACCCGGGGTTAANCCCCGGGC
TTTCACATCCNACNTGACAAGCCGCTACAANCTCTTTACGCCCAATAATTCCGGACAACGCTTGCNCCCTACTT
ATTACCGCGGCTGCTGGCACTTATTTAGCCGGCGCTTCTTCTGCAGGTACCGTCACTTTTCGCTTCTTCCCTGCTN
AAAAAGGTTTACAACCCNAAGGCCGTCATCCCTCAGCGGGCNTCGCTGCATCAGGCTTTCNCCCATTGTGCAATA
TTCCCCACTGCTGCCTCCCGTAGGATTCTGGGCCGNTCTCATTCCANTGTGGCCGGTGCCTCTCAGGCCGG
CTACCCGTCNTCNCCTTGGTAGGCCATTACCCACCAACAAGCTNATAGGCCGCGGGCTCATCCTTCACCGCCGG
AAGCTTTCACCCCNCTCATGCGGGANAATGTTNTCCGGTATTAAACCCCGTTTCCAGGGNTGTCCCAAAT
TGAAGGGGGGATTGNCCACTTTTACTCACCGGTCNCCNCTAATCCACCACC

EN26**SEQ ID NO:11**

CCGCCTTCGCCACCGGTGTTCTCCTGATATCTGCGCATTTACCGCTACACCAGGAATTCCNATCTCCCCTACC
GAACTCTANCTGCCCCGTATCNACTGCAAACCCGGGGTTAAGCCCCGGGCTTTCACCAACCGACNTGACAAGCCGC
CTACAANCTCTTTACNCCCAATAATTCCGGACAACGCTTTCGCCCTACNTATTACCGCGGCTGCTGGCACNTATT
TAGCCGGCGCTTCTTCTGCAGGTACCGTCACTTTTCGCTTCTTCCCTGCTGAAAAAGGTTTACAACCCGAAGGCCG
TCNTCCCTCAGCGGGCTGCTGCATCAGGCTTTCGCCCATTGTGCAATATTCCCACTGCTGCCTCCCGTAGGA
TTCTGGGCCGTGTCTCANTCCANTNTGGCCGGTCCCTCTCAGGCCGNTACCGTCTGCTCCCTGGGTGAACNC
TACCTCNCCAACAANCTGATAGGGCGCGGGCTCANCNTGCACGCCGGANCTTT

EN35**SEQ ID NO:14**

AACACATGCAAGTCGAACGATGAAGCCGCTTCGGTGGTGGATTAGTGGCGAaCGGGTGAgtAACACGtGGCCAAN
TGTGNCCGTCActaTGGGACgAAGaCCTTGGAACGGGGTCTAATACCGGATAACACTCTGTCCCGCATGGGACG
GGTTGAAAGCTCCGGCGGTGAAGGATGAGCCCGCGGCTATCAGCTTGTGGTGGGGTAATGGCCTACCAAGGC
GACGACGGGTAGCCGGCTGAGAGGGCGACCGGCCACACTGGGACTGAGACACGGCCAGACTCCTACGGGAGGC
AGCAGTGGGGAATATTGCACAATGGGCGAAAGCCTGATGCAGCGACGCCGCTGAGGGATGACGGCCTTCGGGTT
GTAAACCTCTTTACGAGGGAAGAAGCGAAAGTGACGGTACCTGCAGAAGAAGCGCCGGCTAACTACGTGCCAGC
AGCCGCGGTAATACGTAGGGCGCAAGCGTTGTCCGGAATTATTGGGCGTAAAGAGCTCGTAGGCGGCTTGTACG
TCGGATGTGAAAGCCCCGGGGCTTAACCCCGGGTCTGCATTGCATACGGGCTAGCTAGAGTGTGGTAGGGGAGATC
GGAATTCTTGGTGTAGCGGTGAAATGCGCAGATATTAGGAGGAACACCGGTGGCGAAGGCGGATCTCTGGGCCA
TTACTGACGCTGAGGAGCGAAAGCGTGGGGAGCGAACAGGATTATATACCCTGGTAGTCCACGCCGTAAACGTTG
GGAAGTGGTGTGGCGACATTCCACGTCGTGCGTCCGCGAGCTAACGCATTAAAGTTCCTCCGCTGGGGAGTACG
CCGCAAGGCTAAAACCTCAAAGGAATTGACGGGGGCCGCACAAGCAGCGGAGCATGTGGCTTAATTCGACGCAA
CGCGAAGAACCTTACCAAGGCTTGACATATACCGGAAGCATCAgAGATGGTGGCCCCCTTGTGGTGGTATACA

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[FIGURE 16 CONTINUED]

GGTGGTGTCATGGCTGTCGTCANCTCGTGTCTGTCGTCGATGTTGGGTTAAGTCCCGCAACGAGCGCAACCCCTTGTTCCT
GTGTTGCCAGCATGCCCTTCGGGGTGATGGGGACTCACAGGAGACTGCCGgGGTCAaCTCGGAGGAAGGTGGGGA
CGACgTCaAGTCATCATGCCCTTATGTCTTGGGCTGCACACGTGCTACAATGGCCGCTACAATGACCTGCgATG
CCGCGgAGGCgGACCGAATCTCAaACAAGCCCGTCTCATTCCGATTGCGGTCTGCaActcCGACCCCATgAAgTCC
GACTtGCTAgTACTCGCACgTCAACaTtGCTGCGCTGaATACgTCCCCGGGCCTTGTACACACCGCCCGTCACGT
CACGAAAGTCGGTAACACCCGAAGCCGGTGGNCCAACCCCTTGTGGGAGGGAGCTGTGCGAA

EN39

SEQ ID NO:15

ccgccttcgccaccggtgttcctcctgatatactgcgcatttcaccgctacaccaggaattccnatctccccctacc
aacctctagctancccggtatcnaatgcaaaccggggttaacccccgggtttcacacccnactnacaanccgc
ctacaaactctttacgccaataattccggacaacgcttgccgctacttattaccgcggtgctggcacttatt
tagccggcgcttcttctgcaggtaccgtcactttcgttcttccctgctgaaaaaggtttacaacccgaaggcng
tcatccctcacgcggtcgtgcatcaggctttcgccattgtgcaatattccccactgctgcctcccgtagna
ntctggggcggtntctcantcccagtggtgncgggtcgccctctcaggccgggtacccgctgctcncctnggtnaacc
attanntcaccaacaagctgataggccgcggtcatccttcaccgcggagcttttaacccctgcccataaaaa
cagangtnttatccggtattanaacccggtttccagg

EN57

SEQ ID NO:17

GTGCTTAACACATGCAAGTCGAACGATGAAGCCGCTTCGGTGGTGGATTAGTGGCGAACGGGTGAGTAACACGTG
GGCAATCTGCCCTTCACTCTGGGACAAGCCCTGGAAACGGGGTCTAATACCGGATAACACTCTGTCCCGCATGGG
ACGGGGTTGAAAGCTCCGGCGGTGAAGGATGAGCCCGCGGCCCTATCAGCTTGTGGTGGGTAATGGCCTACCAAG
GCGACGACGGGTAGCCGGCCTGAGAGGGCGACCGGCCACACTGGGACTGAGACACGGCCAGACTCCTACGGGAG
GCAGCAGTGGGGAATATTGCACAATGGGCGAAAGCCTGATGCAGCGACGCCGCGTGAGGGATGACGGCCTTCGGG
TTGTAAACCTCTTTCAGCAGGGAAGAAGCGAAAGTGACGGTACCTGCAGAAGAAGCGCCGGCTAACTACGTGCCA
GCAGCCGCGGTAAATACGTAGGGCGCAAGCGTTGTCCGGAATTATTGGGCGTAAAGAGCTCGTAGGCGGCTTGTCA
CGTCGGATGTGAAAGCCCGGGGCTTAACCCCGGGTCTGCATTTCGATACGGGCTAGCTAGAGTGTGGTAGGGGAGA
TCGGAATTCTTGGTGTAGCGGTGAAATGCGCAGATATNCAGGAGGAACACCGGTGGCGAAGGCGGATCTCTGGCC
ATTACTGACGCTGAGGAGCGAAAGCGTGGGGAGCGAACAGGATTAGATACCCCTGGTAGTCCACGCCGTAAACGTT
GGGAAC TAGGTGTTGGCGACATTCCACGTCTGTCGGTGCCGCAGCTgAACGCATTAAGTTCCCCGCCTGGGGAGTA
CGGCCGCAAGGCTAAAACCTCAAAGGAATTGACGGGGGCCCCGACAAAGCAGCGGAGCATGTGGCTTAATTCGACGC
AACGCGAAGAACCTTACCAAGGCTTGACATATACCGGAAAGCATCAGAGATGGTGCCCCCTTGTGGTGGGTATA
CAGGTGGTGCATGGCTGTCTGTCAGCTCGTGTCTGTCGTCGATGTTGGGTTAAGTCCCGCAACGAGCGCAACCCCTTGT
CTGTGTTGCCAGCATGCCCTTCGGGGTGATGGGGACTCACAGGAgACTGcCGGGGTCAACTCGGAGGAAGGTGGG
GACGACGTCAAGTCATCATGCCCTTATGTCTTGGGCTGCACACGTGCTACAATGGCCGGTACAATGAGCTGCGA
TGCCgCGAgGCGGAgCgAATCTCAAAAAGCCGGTCTCAGTTCGGATTGGGGTCTGCAACTCGACCCCATGAAGTC
GGAgTTGCTAgTAATCgCAGATCAGCATTGCTGcGGTGAATACGTTCCCGGGCCTTGTACACACCGCCGTCAcGT
CACGAAAGTCGGTaACACCCGAAGCCGGTGGCCCAACCgCCTTGTGGGAgGGAACCTTTCCA

FIGURE 17

SE1

SEQ ID NO:19

GAACGATGAAGCCGTTTTCGGTGGTGGATTAGTGGCGAACGGTGAGTAAAAGTGGCAATTTNCCCTTCATTTTGGAC
CAAGCCCTGGAAACGGGTTTAAANACCGGATAACATTTNTGTCCCGCATGGGACGGGGTTGAAAGNTCCCGGCGGTG
AAGGATGAGCCCGCGGCNTATCAGCTTGTGGTGGGGTAATGGCCTACCAAGGCGACGACGGGTAGCCGGCCTGA
GAGGCGACCGGCCACACTGGGANTGAGACACGGCCAGACTCCTACGGGAGGCAGCAGTGGGGAATATTGCACA
ATGGGCGAAAGCCTGATGCAGCGACGCCGCGTGAGGGATGACGGCCTTCGGGTTGTAAACCTNTTTCAGCAGGGA
AGAAGCGAAAGTGACGGTACCTGCAGAAGAAGCGCCGGCTAAATANGTGCCAGCAGCCGCGGTAATANGTAGGGC
GCAAGCGTTGTCCGGAATTATTGGGCGTAAAGAGCTTGTAGGCGGCTTGTGANGTCGGATGTGAAAACCCGGGGC
TTAACCCCGGGTTTGCATTTCGATACGGGCTAGTTAGAGTGTGGTAGGGGAGATNGGAATTCCTGGTGTAGCGGTG
AAATGCGCAGATATCAGGAGGAACACCGGTGGCGAAGGCGGATCTCTGGGCCATTACTGACGCTGAGGAGCGAAA
GCGTGGGGAGCNAACAGGATTAGATACCTGGTAGTCCACGCCGTAAACGTTGGGAAGTAGGTGTGGCGACATT
CCACGTCGTGGTGCCGAGCTAACGCATTAAGTTCCCGCCTGGGGAGTAGCGCCGCAAGGCTAAAACCTCAAAG
GAATTGACGGGGGGCCCGACAAGCAGCGGAGCATGTGGCTTAATTGACGCAACGCGAACAACCTTACCAAGGCT
TGACATATACCGGAAAGCATCANAGATGGTGCCCCCTTGTGGTTCGGTATACANGTGGTGCATGGCTGTCGTGAG
CTCGTGTGTCGTGAGATGTTGGGTTANGTCCCGCAACGAGCGCNACCCTTGTTCGTGTCGNCNAGCATGCCCTTCG
NGGTGATGGGGACTCACANGAGACTGNCGGGGTCCACTCGGAGGAAGGTGGCGACNACGTCANNTCATCATGCC
CCTTATGTCTTGGNCTGGCCACGTGCNACNATGGCC

SE2

SEQ ID NO:20

GCTGGCGCGTGCTTAACACATGCAAGTCGAACGATGAAGCCGCTTCGGTGGTGGATTAGTGGCGAACGGGTGAG
TAACACGTGGGCAATCTGCCCTTCACTCTGGGACAAGCCCTGGAAACGGGGTCTAATACCGGATAACACTCTGTC
CCGCATGGGACGGGGTTGAAAGCTCCGGCGGTGAAGGATGAGCCCGCGGCCTATCAGCTTGTGGTGGGGTAATG
GCCTACCAAGGCGACGACGGGTAGCCGGCCTGAGAGGGCGACCGGCCACACTGGGACTGAGACACGGCCAGACT
CCTACGGGAGGCAGCAGTGGGGAATATTGCACAATGGGCGAAAGCCTGATGCAGCGACGCGCGTGAGGGATGAC
GGCCTTCGGGTTGTAAACCTCTTTCAGCAGGGAAGAAGCGAAAGTGACGGTACCTGCAGAAGAAGCGCCGGCTAA
CTACGTGCCAGCAGCCGCGGTAATACGTAGGGCGCAAGCGTTGTCCGGAATTATTGGGCGTAAAGAGCTCGTA_g
CGGCTTGTACGTCGGATGTGAAAGCCCGGGGCTTAACCCCGGGTCTGCATTTCGATACGGGCTA_gCTA_gAGTGTG
GTAGGGGAGATCGGAATTCCTGGTGTAGCGGTGAAATGCGCaGATATCA_gGAGGAACaCGGTGGg_gAAGgcGGA
TCTCTGGGcCaTACTGACGCTGAGGAGCGAAAGCGTGGGGAGCGAACAGGATTAGATACCTGGTAGTCCAAGC
CGTAAACGTTGGGAACANGTGTGGCGACATTCCACGTCGTGGTGCCGAGCTAACGCATTAAGTTCCCGGTC
CTGGGGAGTACGGCCGCNAGGCTAAAACCTCAAAGGAATTGACGGGGGGCCCGCACAAGCAGCGGAGCATGTGGCTT
ANTTCGACGCNACGCGAAGAACCTTNCCAAGGCTGACATATACCGgAAAGCATCaAGATGGTGCCCCCTTGTG
GTCGGTATACAGGgTGGTGCATGGCTGTtCGtCaGCTCGTGTCgtGAGATGTTGGGTTAaGTCCCGCAAAGAGCG
CAACCgTGTTCGTGTGTGCCAGCATGCCCTTCGGGGTGATGGGGACTCaCaGAgACTGTCNGGGTCAACTCgga
GGAAGGTGGgGACGACgTCAAGTtCATCATGCCCTTATGTCTTGGGCTGCACACNGCTACAATGGCCGGTACA
ATGAGNNGGGATGCCGCGAGGCGGAGCGAATCTCAAAAAGCCGGTCTCAGTTCGGATTGGGGTCTGCAACTGACC
CCATGAAGTCGGAGTTGCTAGTAATCGCAGATCAGCATTGCTGCGGTGAATACGTNCCCGGGCCTNGTACACACC
ACCCGTCACGTACGAAAGTCGGTAACACCCTAAGCCGGTGNCCCAACCCCTTNTGGGAGG

FIGURE 18

PM36

SEQ ID NO:21

CCAGANATCCGCCTTCGCCACCGGTGTTCCCTCCTGATATCTGCGCATTTACCGCTACACCAGGAATTCCGATCT
CCCCTACCACACTCTAGCTAGCCCGTATCGAATGCAGACCCGGGGTTAAGCCCCGGGCTTTCACATCCGACGTGA
CAAGCCGCCTACGAGCTCTTTACGCCCAATAATTCGGGACAACGCTTGCGCCCTACGTATTACCGCGGCTGCTGG
CACGTAGTTAGCCGGCGCTTCTTCTGCAGGTACCGTCACTTTCGCTTCTTCCCTGCTGAAAGAGGTTTACAACCC
GAAGGNCGTATCCCTCACGCGGCGTCTGCTGCATCAGGCTTTCGCCCATTTGTGCAATATTCCCCACTGCTGCCTC
CCGTAGGAGTCTGGGNCGTGTTCAATNCCAGTGGTGGGCGGCTCGCCCTCTCAGGNCGGCTACCGTCTCGCCTT
GGTAGGCATTACCACAACAAGCTGATAGGCGGGGGTTCATCCTTCAACGCCGGAGCTTCAAACCCGTCCATGCGGG
ACAAGTGTATCCGGTATTAAACCC

PM40

SEQ ID NO:22

TCAGTNATGGCCCAGAANGATCCGNCTTCGCCACCGGTGTTCCCTCCTGATATCTGCGCATTTACCGCTACACCA
GGAATTCGGATCTCCCCTACCACACTCTAACTAGCCCGTATCGAATGCAGACCCGGGGTTAAGCCCCGGGCTTTC
ACATCCGACGTGACAAGCCGCCCTACGAGCTCTTNACGCCCAATAATTCGGGACAACGCTTGCGCCCTACGTATTA
CCGCGGCTGCTGGCACGTAGTTAGCCGGCGCTTCTTCTGCAGGTACCGTNACTTTCGCTTCTTCCCTGCTGAAAG
AGGTTTACAACCCGAAGGCCGTCNTCCCTCACGCGGCGTCTGCTGCATCAGGCTTTCGCCCATNGTGCANTATTCC
CCACTGNTGNTCCCCTANGAGTCTGGGCGGTGTCTCAGTCCCAGTGTGGCCGGTCGNCTCTCAGGCCGGCTAC
CGTCGTCGCCTTGGTAGGNCATTACCCACCAACAAGCTGATANGTCGNGGGCTCATCCTTACCGNCGGAGNTTT
AACCCCGTNCATGCGGGACAGAGTGTTATCCGGTATTANACCCGTATNCAGGGCTTGTCCCATAGTGAAGGGNAG
ATNGCCACGTGTTATCACCCTTCGNCACTAATNATCANCGAANC GGCTTCATCGTTCGACTTG CATGTGTTA

PM41

SEQ ID NO:23

CTCAGCGTCAGTCATGGCCAAGAGATCCGCCTTCGCCACCGGTGTTCCCTCCTGTATATCTGCGCATTTACCGCT
ACACCAGGAATTCGGATCTCCCCTACCACACTCTAGCTAGCCCGTATCGAATGCAGACCCGGGGTTAAGCCCCGG
GCTTTCACATCCGACGTGACAAGCCGCCCTACGAGCTCTTTACGCCCAATAATTCGGGACAACGCTTGCGCCCTAC
GTATTACCGCGGCTGCTGGCACGTAGTTAGCCGGCGCTTCTTCTGCAGGTACCGTCACTTTCGCTTCTTCCCTGC
TGAAAGAGGTTTACAACCCGAAGGCCGTCATCCCTCACGCGGCGTCTGCTGCATCAGGCTTTCGCCCATTGTGCAA
TATTCCCCACTGCTGCCTCCCCTAGGAGTCTGGGCGGTGTCTCAGTCCCAGTGTGGCCGGTCGCCCTCTCAGGCC
GGCTACCCGTCGTCGCCTTGGTAGGCCATTACCCACCAACAAGCTGATAGGCCGCGGGCTCATCCTTCANCGNCG
GAGCTTTAACCCGTCATGCGGGACAGAGTGTTATCCGGTATTAAACCCGTTTCAGGGCTTGTCCCANAGTGAAG
GGCAGATTGCCACGTGTTATCANCCGTTTCGNCACTAATCACANCGAANC GGTTTCATCGTTCGACTTG CATGTGT
TAA

PM171

SEQ ID NO:25

CCCTCAGGGTCAGTAATGGGCCCAGAGATCCGCCTTCGCCACCGGTGTTCCCTCCTGAATATCTGCGCATTTACCC
GCTACACCAGGAATTCGGATCTCCCCTACCACACTCTAGCTAGCCCGTATCGAATGCAGACCCGGGGTTAAGCCC
CGGGCTTTCACATCCGACGTGACAAGCCGCCCTACGAGCTCTTTACGCCCAATAATTCGGGACAACGCTTGCGCCC
TACGTATTACCGCGGCTGCTGGCACGTAGTTAGCCGGCGCTTCTTCTGCAGGTACCGTCACTTTCGCTTCTTCCC
TGCTGAAAGAGGTTTACAACCCGAAGGCCGTCATCCCTCACGCGGCGTCTGCTGCATCAGGCTTTCGCCCATTGTG
CAATATTCCCCACTGCTGCCTCCCCTAGGAGTCTGGGCGGTGTCTCAGTCCCAGTGTGGCCGGTCGCCCTCTCAG
GCCGGCTANCCGTCGTCGCCTTGGGTAGGCATTANCCCANCAACAAGCTGATAGGNCGCGGGCTCATNCTTCAAC
GCCGGAGCTTTCAANCCCGTCCATGCGGGACAGAGTGTTATNCGGTATTAAACCCGTTTCAGGGCTTGTTCAGAG
GTGAAGGGCAGATTGCCACGTGTTATCAACCGTTTCGGCACTAATCACACGAAGCGGNTTATCGTTCGACTTGCA
TGTGTTAAACAAGCCGCCAGCGTTCGTC

[FIGURE 18 CONTINUED]

PM185

SEQ ID NO:26

TCAGTAATGGCCCAGAGATCCGCCTTCGCCACCGGTGTTCTCCTGGATATCTGCGCATTTACCGCTACACCAG
GAATTCCGATCTCCCTACCACACTCTAGCTAGCCCGTATCGAATGCAGACCCGGGGTTAAGCCCCGGGCTTTCA
CATCCGACGTGACAAGCCGCCTACGAGCTCTTTACGCCCAATAATTCCGGACAACGCTTGCGCCCTACGTATTAC
CGCGGCTGCTGGCACGTAGTTAGCCGGCGCTTCTTCTGCAGGTACCGTCACTTTGCTTCTTCCCTGCTGAAAGA
GGTTTACAACCCGAAGGCCGTATCCCTCACGCGCGCTCGCTGCATCAGGCTTTGCCCCATTGTGCAATATTCCC
CACTGCTGCCTCCCGTAGGAGTCTGGGCGGTGTCTCAGTCCCAGTGTGGCCGGTTCGCCCTCTCAGGCGCGGTACC
CGTCGTCGCCTTGGTAGGCCATTACCCACCAACAAGCTGATAGGCCGCGGGCTCATCCTTCACCGNCGGAGCTT
TAACCCCGTCCCATGCGGGACAGAGTGTTATCCGGTATTAGAACCCGTTTCCAGGGCTTGTCCCAGAGTGAAGGG
CAGATTGCCACGTGTTACTCANCCGTTTCGNACTAATCANCAACGAAGCGGCTTCATCGTTCGACTTGCATGTGT
TAAGCACGCCGNCAGCGTTCGTCTGAGCCAGGATC

PM208

SEQ ID NO:27

TCAGTATCNGCCCAGAGATCCGCCTTCGCCACCGGTGTTTCTCCTGATATCTGCGCATTTACCGCTACACCAG
GAATTCCGATCTCCCTACCGAACCTCTAGCCTGCCCGTATCGACTGCAGACCCGGGGTTAAGCCCCGGGCTTTCA
CAACCGACGTGACAAGCCGCCTACGAGCTCTTTACGCCCAATAATTCCGGACAACGCTTGCGCCCTACGTATTAC
CGCGGCTGCTGGCACGTAGTTAGCCGGCGCTTCTTCTGCAGGTACCGTCACTTTGCTTCTTCCCTGCTGAAAGA
GGTTTACAACCCGAAGGCCGTATCCCTCACGCGCGCTCGCTGCATCAGGCTTTGCCCCATTGTGCAATATTCCC
CACTGGTGNCTCCCGTANGAGTCTGGGGCGGTGTCTCANTCCAGTGTGGGCGGTTCGCCCTCTCAGGGCGGCTACCGT
CGTCGCTTGGTGAGNCACTACTCACAACAAGCTGATAGGCCGCGGGCTCATCTGGAACGGCGGAGCTTTACAC

PM228

SEQ ID NO:28

TCAGTAATGGCCCAGANATCCGNCTTCGCCACCGGTGTTCTCCTGATATCTGCGCATTTACCGCTACACCAGG
AATTCCGATCTCCCTACCACACTCTAACTAGCCCGTATCGAATGCAGACCCGGGGTTAAGCCCCGGGCTTTAC
ATCCGACGTGACAAGCCGCCTACGAGCTCTTTACGCCCAATAATTCCGGACAACGCTTGCGCCCTACGTATTACC
GCGGCTGCTGGCACGTAGTTAGCCGGCGCTTCTTCTGCAGGTACCGTCACTTTGCTTCTTCCCTGCTGAAAGAG
GTTTACAACCCGAAGGCCGTATCCCTCACGCGCGCTCGCTGCATCAGGCTTTGCCCCATTGTGCAATATTCCCC
ACTGCTGCCTCCCGTANGAGTCTGGGCGGTGTCTCAGTCCCAGTGTGGCCGGTTCGCCCTCTCAGGCCGGCTACCG
TCGTGCGCTTGGTAGGCCATTACCCACCAACAAGCTGATANGNCGNGGGCTCATCCTTCACCGNCGGAGCTTTCA
ANCCCGTCCCATGCGGGACAGAGTGTTATCCGGTATTAAACCCGNTTCCAGGGCTTGTCCATAGTGAAGGGCAGA
TTGCCAAGTGTATCANCCGTTTCGNACTAATCATCANCAAGCGGCTTCATCGTTCGACTGCATGTGTT

PM252

SEQ ID NO:29

TCCTCAGNATCAGTAATGGCCCAGAGATCCGCCTTCGCCACCGGTGTTCTCCTGATATCTGCGCATTTACCGC
TACACCAGGAATTCCGATCTCCCTACCACACTCTANCTAGCCCGTATCGAATGCAGACCCGGGGTTAAGCCCCG
GGCTTTCACATCCGANGTGACAAGCCGCCTACGAGCTCTTTACGCCCAATAATTCCGGACAANGCTTGCGCCCTA
CGTATTACCGCGGNTGCTGGCACGTAGTTAGCCGGCGCTTCTTCTGCAGGTACCGTCACTTTGCTTCTTCCCTG
CTGAAAGAGGTTTACAACCCGAAGGCCGTATCCCTCACNCGGCGTTCGCTGCATCAGGCTTTGCCCCATTGTGCA
ATATTCCCCACTGCTGCCTCCCGTAGGAGTCTGGGCGGTGTCTCAATCCCANTGTGGCCGGTTCGCCCTCTCANGC
CGGCTACCGTTCGTCGCTTGGTAGGCCATTACCCACCAACAAGCTGGATAGGNCGGGGGCTCATCTTTCACCGCC
GGAAGCTTTAANCCCGTCCATGCGGGANANAGTGNATCCCNATTAACCCNGTTTCAGGGCTTGTCCANAGTG
AAGGNGATTGCCNAGTGTATTCNCCCGTTCGCCANTAATCNACAACGAAGCGGNTTCNCTCGNTTCGACTTG
C

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[FIGURE 18 CONTINUED]

PM342

SEQ ID NO: 30

TAATGGCCCAGAAATCCGCCTTCGCCACCGGTGTTCCCTCCTGAATATCTGCGCATTTACCGCTACACCAGGAA
TTCCGATCTCCCCCTACCACACTCTAGCTAGCCCGTATCGAATGCAGACCCGGGGTTAAGCCCCGGGCTTTCACAT
CCGACGTGACAAGCCGCCTACGAGCTCTTTACGCCCAATAATTCCGGACAACGCTTGCGCCCTACGTATTACCGC
GGCTGCTGGCACGTAGTTAGCCGGCGCTTCTTCTGCGAGGTACCGTCACTTTCGCTTCTTCCCTGCTGAAAGAGGT
TTACAACCCGAAGGCCGTATCCCTCACGCGGCGTCGCTGCATCAGGCTTTCGCCCATTGTGCAATATTCCCCAC
TGCTGCCTCCCGTAGGAGTCTGGGCGGTGTCTCAGTCCCAGTGTGGCGGTGCGCCCTCTCAGGCCGGNTANCCGTC
GTCGCCTTGGTANGCCATTANCCCAACAAGCTGATANGCCGNGGGCTCATCCTTCANCGCCGGAGCTTTTAA
CCCCGTCCCATGCGGGACAGAGTGTTATCCGGTATTAGATCCCGTNTCCAGGGCTTGTNCATAGTGAAGGGCANA
TTGCCACGTGTTACTCANCCGTTCGC

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FIGURE 19

EN4

GGCGGCGTGCTTAACACATGCAAGTCGAACGATGAACCACTTCGGTGGGGATTAGTGGCGAACGGGTGAGTAACA
CGTGGGCAATCTGCCCTTCACTCTGGGACAAGCCCTGGAAACGGGGTCTtAATACCGGATAACACTCCCCTCTC
CTGAGTGGGGGTAAAAGCTCCGGCGGTGAAGGATGAGCCCGCGGCCTATCAGCTTGTGGTGAGGTAATGGCTC
ACCAAGGCGACGACGGGTAGCCGGCCTGAGAGGGCGACCGGCCACACTGGGACTGAGACACGGCCCCAGACTCCTA
CGGGAGGCAGCAGTGGGGAATATTGCACAATGGGCGAAAGCCTGATGCAGCGACGCCCGGTGAGGGATGACGGCC
TTCGGGTTGTAAACCTCTTTCAGCAGGGAAGAAGCGAAAGTGACGGTACCTGCAGAAGAAGCGCCGGCTAACTAC
GTGCCAGCAGCCCGGTAATACGTAGGGCGCAAGCGTTGTCCGGAATTATTGGGCGTAAAGAGCTCGTAGGCGGC
TTGTACAGTCGGGTGTGAAAGCCCGGGGCTTAACCCCGGTCTGCATTGATAACGGGCTAGCTAGAGTGTGGTAG
GGGAGATCGGAATTCCTGGTGTAGCGGTGAAATGCGCAGATATTGAGGAGGAACACCGGTGGCGAAGGCGGATCT
CTGGGCCATTACTGACGCTGAGGAGCGAAAGCGTGGGGAGCGAACAGGATTAGATACCCTGGTAGTCCACGCCGT
AAACGGTGGGAACTAGGTtGTTGGCGACATTCCACGTGCTCGGTGCCGCGAGCTAACGCATTAAGTTCCCCGCGCTG
GGGAGTACGGCCGCAAGGCTAAACTCAAAGGAATTGACGGGGGGCCCGCACAAAGCAGCGGAGCATGTGGCTTAAT
TCGACGCAACGCGAAGAACCCTTACCAAGGCTTGACATACACCGGAAAGCATCAGAGATGGTGCCCCCTTGTGGT
CGGTGTACAGGTGGTGCATGGCTGTCTGCTCAGCTCGTGTCTGAGATGTTGGGTAAAGTCCCGCAACGAGCGCAAC
CCTTGTCTGTGTGTCAGCATGCCCTTCGGGGTGATGGGGACTCACAGGAGACCGCCGGGTCAACTCGGAGGAA
GGTGGGGACgACGTCAAGTCATCATGCCCTTATGTCTTGGGCTGCACACGTGCTACAATGGCAGGTACAATGAG
CTGCGATACCGTGAGGTGGAGCGAATCTCAAAAAGCCTGTCTCAgTTCGGATTGGGGTCTGCAACTCGACCCcaT
GAAGTCGGAGTTGCTAGTAATCGCAGATCAGCATTGCTGCGGTGAATACGTTCCCGGGCCTTGTACACACCGCCC
GTCACGTACGAAAGTCGGTAACACCCGAAGCCGGTGGCCCAACCCCTTGTGGGAGGGAGCTGTCAAGGTGGGA
CTGGCGATTGGG

EN10

GAGTTTGATCANNGGCTCAGACGAACGCTGGCGGCGTGTTAACACAANCCAAGTCGAANGNTGAACCACTTCGTTG
GGATTAGTGCGAACGGTGNTAACACGNTGGCAATGTGCCCTTCACTNTGGGACAAGNCCTGGAACCGGGTTCTA
ATACCGGATAACCACTACCCGACGGCATCTGTGGTGTGTTGAAAGCTCCGCCGTGAAGGATGAGCCCGCGGCCTAT
CAGCTTGTGGTGAGGTAATGGCTCACCCAAGGCGACGACGGATAGCCGGCCTGAGAGGGCGACCGGCCACACTG
GGACTGAGACACGGCCAGACTCCTACGGAGGCGACGAGTGGGGAATATTGCACAATGGGCGAAAGCCTGATGCAG
CGACGCCGCGTGAGGGATGACGGCCTTCGGGTTGTAAACCTCTTTCAGCAGGGAAGAAGCGAAAGTGACGGTACC
TGCAGAAGAAGCGCCGGCTAACTACGTGCCAGCAGCCGCGGTAATACGTAGGGCGCAAGCGTTGTCCGGAATTAT
TGGGCGTAAAGAGCTCGTAGGCGGCTTGTACGTCGGGTGTGAAAGCCCGGGGCTTAACCCCGGTCTGCATTG
ATACGGGCTAGCTAGAGTGTGGTAGGGGAGATCGGAATTCCTGGTGTACCCGGTGAAATGCGCAGATATCAGGAG
GAACACCGGTGGCGAAGGCGGATCTCTGGGCCATTACTGACGCTGAGGAGCGAAAGCGTGGGGAGCGAACAGGAT
TAGATACCCTGGTAGTCCACGCCGTAAACGGTGGGAACTAGGTGTTGGCGACATTCCACGTCGTGGTGCCGCGAG
CTAACGCATTAAGTTCCCCGCTGGGGAGTACGGCCGCAAGGCTAAACTCAAAGGAATTGACGGGGGCCCGCAC
AAGCAGCGGAGCATGTGGCTTAATTGACGCAACGCGAAGAACCCTTACCAAGGCTTGACATACGCCGGAAGCAT
CAGAGATGGTGCCCCCTTGTGGTGGTGTACAGGTGGTGCATGGCTGTCTGCTCAGCTCGTGTCTGAGATGTTGG
GTTAAGTCCCGCAACGAGCGCAAcCCTTGTCTGTGTTGcCAGCATGCCCTTCGGGGTGATGGGGACTCACAGGA
GACCGCCGGGGTCAACTCGGAGGAAGGTGGNGACGACGTCgAGTCATCATGCCCTTATGTCTTGGGGCTGCACA
CGTGCTACNATGggCaGGTACAATGAGCTGCGATACCGTGAGGTGGAGCgCATCTnnnnnAGCctGTCTCAGTTC
GgATTGGGGTCTGcAACTCGACCCCaTGAAGTCGgAGTTGCTAgATAATCgCAgATCAGCATTGctGCGgTGAAT
ACGTtCCCCGGCCTTGTACACACCGCCCGTCACGTACGAAAGTCGGTAACACCCGAAGCCGGTGGCCCAACCC
TTGTGGGAGGGAGCTGTCAANGTGG

[FIGURE 19 CONTINUED]**EN22**

TCCTCAGCGTCAGTAATGGCCCCAAAAACCGCCTTCGCCACCGGTGTTCTCTGATATCTGCGCATTTACCGCT
ACACCAGGAATTCCNATCTCCCCTACCACACTCTAGCTAGCCCGTATCNAATGCAAACCCGGGGTTAANCCCCGG
GCTTTACATCCNACNTGACAAGCCGCTTACAANCTCTTTACGCCCAATAATCCGGACAACNCTTGCGCCCTAC
TTATTACCGCGGCTGCTGGCACTTATTTAGCCGGCGCTTCTTCTGCAGGTACCGTCACCTTTCGCTTCTTCCCTGC
TGAAAAAGGTTTACAACCCNAAGGCCGTCATCCCTCACGCGGNTCNCCTGCATCAGGCTTTCNCCCATTGTGCAA
TATTCCTCCACTGCTGCCTCCCGTAGGATTCTGGGCGTNTCTCANTCCANTGTGGCCGGTCGCCCTCTCAGGCC
GGCTACCCGTCGTCNCCTTGGTAGGCCATTACCCCNCCAACAANCTGATAGGCCGCGGGCTCNTCTTCACCGCC
GGAGCTTTCAACCCCGTCCCATGCGGGANAAANTGTTNTCCGGTATTAAAACCCGTTTCCAGGGNTTGTCCAAA
TTGAAGGGNANATTGCCCACTTTTNNTCACCCGTTCCCACTAATCCACCACCGAA

EN30

TGGNGNGTGCTTAACACATGCAAGTCGAACGATGAANCCTTTGGGGTGGATTAGTGGCGAACGGGTGAGTAAC
ACGTGGGCAATCTGCCCTTCACTCTGGGACAAGCCCTGGAACGGGGTCTAATACCGGATAACACTCTGTCCCGC
ATGGGACGGGGTTAAAAGCTCCGGCGGTGAAGGATGAGCCGCGGCCTATCAGCTTGTGGTGGGGTGATGGCCT
ACCAAGGCGACGACGGGTAGCCGGCTGAGAGGGCGACCGCCACACTGGGACTGAGACACGGCCAGACTCCTA
CGGGAGGCAGCAGTGGGGAATATTGCACAATGGGCGAAAGCCTGATGCAGCGACCCGCGTGAGGGATGACGGCC
TTCGGGTGTAAACCTCTTTCAGCAGGGAAGAAGCGAAAGTGACGGTACCTGCAGAAGAAGCGCCGGCTAACTAC
GTGCCAGCAGCCGCGGTAATACGTAGGGCGCAAGCGTTGTCCGGAATTATTGGGCGTAAAGAGCTCGTANGCGGC
TTGTACGTCGGATGTGAAAGCCCGGGGCTTAACCGGGTCTGCATTGATACGGGCTAACTAAAATGTGGTAGG
GGAGATCGGAATTCTTGGTGTANCGGTGAAATGCGCAGATATCAAGAGGAACANCGGTGGCGAANGCGGATCTCT
GGGCCATTACTGACGCTGAGGAGCGAAAGCGTGGGGAGCGAACAGGATTAGATACCTGGTAGTCCACGCCGTAA
ACGTTGGGAAC TAGGTGTTGGCGACATTCCACGTCGTCGGTGCCGAGCTAACGCATTAAGTTCCCGCCTGGGG
AGTACGGCCGCAAGGCTAAAACCTCAAAGGAATTGACGGGGCCCGCACAAAGCAGCGGAGCATGTGGCTTAATTG
ACGCAACGCGAAGAACCTTACCAAGGCTTGACATATACCGAAAGCATCAGAGATGGTGGCCCCCTTGTGGTGG
TATACAGGTGGTGATGGCTGTCTGTCAGCTCGTGTCTGAGATGTTGGGTAAAGTCCCGCAACGAGCGCAACCCCT
TGTTCTGTGTTGCCAGCATGCCCTTCGGGGTGATGGGACTCACAGGAGACTGCCGGGGTCAACTCGGAGGAAGG
TGGGGACGACGTCAAGTCATCATGCCCTTATGTCTTGGGCTGCACACGTGCTACAATGGCCGGTACAATGAGCT
GCGATGCCGCGAGGCGGAGCGAATCTCAAAAAGCCGGTCTCAGTTCGGATTGGGGTCTGCAACTCGACCCcATGA
AGTCGgAGTTGCTAgTAATCgAgATCAgCATTGCTGCGGTGAATACgTTNCCGGGCCTtGTACAcACCGCCGT
CACGTCACgAAAGTCGGTAACACCCGAAGNCCGGTGGTCCAACCCCTTGTGGGAGGGAGACTGTGCAAGGTGGGA
CTGGCGATTGG

EN43

CTCAGCGTCACTATCGGCCCAAAANACCGCCTTCGCCACCGGTGTTCTCTGATATCTGCGCATTTACCGCTA
CACCAGGAAATTCCANTCTCCCCTACCGAACTCTANCTGCCCGTATCAACCGCAGGCTTGGGGTTAAGCCCCAA
TTTTTCACGGTCAACGCNACAAGCCGCTTACAAGCTCTTTACGCCCAATAAATCCGGACAACGCTCGCACCCCTAC
TTCTTACCGCGGCTGCTGGCACTTATTTGGCCGGTGCTTCTTCTGCAGGTACCGTCACTCTCGCTTCGTCCCTGC
TNAAAAGGTTTACAACCCGAAGGCCGTCATCCCTCACGCGGNTCGCTGCATCAGGCTTCCGCCCATTGTGCAA
TATTCCTCCACTGCTGCCTCCCGTAGGATTCTGGGCGTNTCTCANTCCAGTGTGGCCGGTCGCCCTCTCAGGCC
GGCTACCCGTCGTCGCCTTGGTAGGCCATACCCCAACCAAGCTGATAGGCCGCAAGCCCATCCAAGCCGA
AAAACCTTTCCACCACCAGCCATGCGGCCAAAATTCCTATTTCGGTATTAGCCCCCGTTTCCNAAGGTTNTCCCAA
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AT

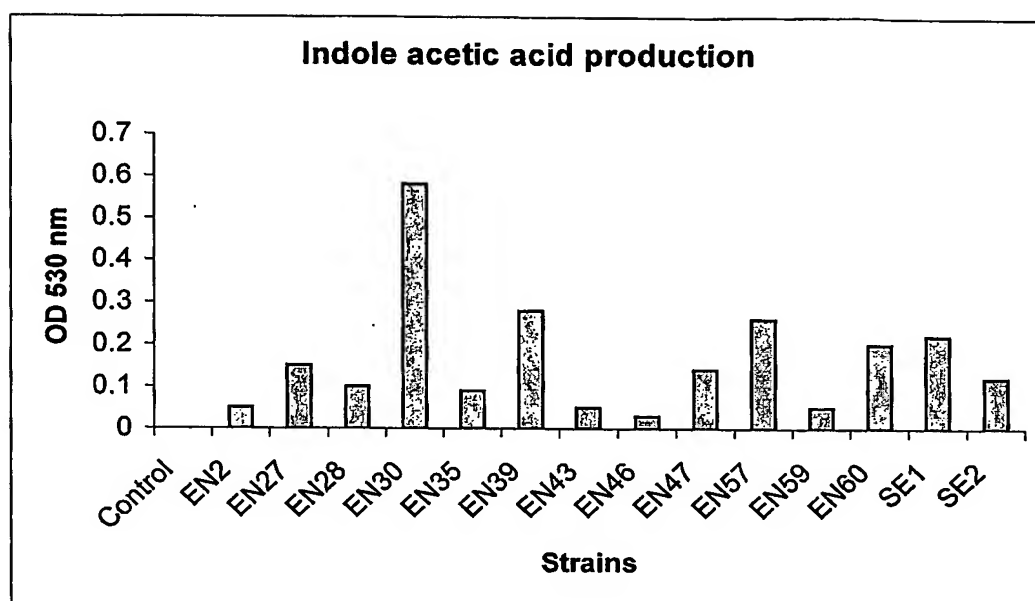
[FIGURE 19 CONTINUED]**EN47**

CGCTGGCGGCGTGCTTAACACATGCAAGTCGAGCGGAAAGGCCCTTCGGGgTACTCGAGCGGGCGAACGGGTGAGT
AACACGTGAGTAATCTGCCCCAGGCTCTGGGATAGCCACCGGAAACGGTGATTAATACCGGATACGACAACCGAT
TGCATGATCTGGTTGTGGAAAGTTTTTCGGCCTGGGATGTGCTCGCGGCCCTATCAGCTTGTGGTGAGGTAATGG
CTCACCAAGGCTTCGACGGGTAGCCGGCCTGAGAGGGTGACCGGCCACACTGGGACTGAGACACGGCCCAGACTC
CTACGGGAGGCAGCAGTGGGGAATATTGGACAATGGGCGGAAGCCTGATCCAGCAACGCCGCGTGAGGGATGACG
GCCTTCGGGTTGTAAACCTCTTTCAGCACAGACGAAGCGCAAGTGACGGTATGTGCAGAAGAAGGACCGGCCAAC
TACGTGCCAGCAGCCGCGGTAATACGTAGGGTCCGAGCGTTGTCCGGAATTATTGGGCGTAAAGGGCTCGTAGGC
GGTCTGTGCGGTCGGGAGTGAAAACCAAGTGCTTAACACCTGGCCTGCTTTCGATACGGGCAGACTAGAGGTACT
CAGGGGAGAATGGAATTCCTgGTGTAGCGGTGAAATGCGCAGATATTAGGAGGAACACCGGTGGCGAAGGCGGT
TCTCTGGGAGTATCCTGACGcTTGAAGAGCGAAAGTGTTGGGGAGCGAACAGGATTAGATANTNTGGTAGTCCACA
CCGTAAACgTTGGGCGcTAGgTGTGGGACACATTCCACGTGTTCCGTGCCGCAGCTAACGCATTAAGCGCCCCGC
CTGGGGAGTACGGCCGCAAGGCTAAACTCAAAGGAATTGACGGGGGCCGACAAGCGGCGGAGCATGCGGATT
AATTTCGATGCAACGCGAAGAACCTTACCTGGGTTTGACATACACCGGAAAGCCGTAGAGATACGGCCCCCTTTTAG
TCGGTGTACAGGTGGTGATGGCTGTCGTGCTGAGATGTTGGGTTAAGTCCCACAACGAGCGCAA
CCCTCGTCCTATGTTGCCAGCAATTCGGTTGGGGACTCATAGGAGACTGCCGGGGTCAACTCGGAGGAAGGTGGG
GATGACGTCAAGTCAATGCTGCCCCCTATGTCCAGGGCTTACGCATGCTACAATGGCCGGTACAAAGGGCTGCGA
TCCCGTGAGGGTGAGCGAATCCCAAAAAGCCGGTCTCAGTTCGGATTGGGGTCTGCAACTCGACCCCATGAAGTC
GGAGTCGCTAGTAATCGCAGATCAGCAACGCTGCGGTGAATACGTTCCCGGCCCTTGTACACACCGCCCGTCACG
TCACGAAAGTCGGCAACACCCGAAGCCAGTGGCCCAACCCTTGTGGGGGAGCTGTCAAGGTGGGGCTGGCGAT
TG

EN59

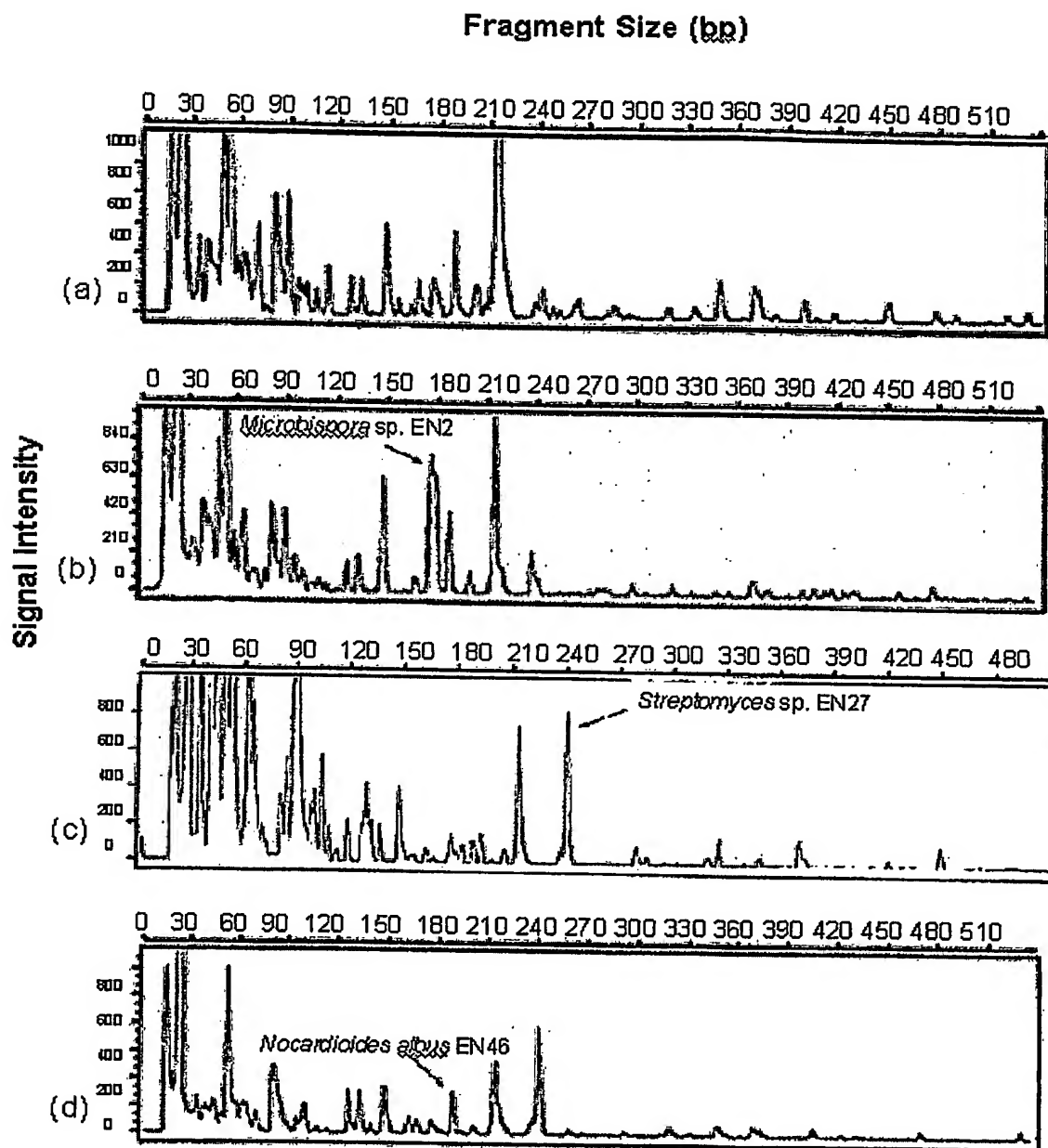
GGGNATTAGTGGGGAACGGGTGAGTAAAANGTGGCCANTTTCCCCTGNATTTTGGACANCCCCNGGAAANGGNTT
NTAAAACNGGATANTGACCACCTTGGCATCCAAGTTTTNGAACTTCGGGCGGTGCAGGATGAGCCNCGGCNTA
TNAGCTTGTTGGNGAGGTAATGGNTCACCAGGGGANGACGGGTAGCCGGCCTGAGAGGGGACCNGCCACANTGGG
ANTGAGANACGGCCCAGANTCCTACGGGAGGCAGCAGTGGGGAATATTGCACAATGGGCGAAAGCCTGATGCAGG
ANNCCGCGTGAGGGANGACGGCCTTNGGGTTGTAAACNTTTTTNAGCAGGGAAGAAGCGAAAGTGACGGTACCTG
CAGAAGAAGCGCCGGCTAAATAAGTGCCAGCAGCCGCGTAATAAGTAGGGNGCGAGCGTTGTNCGGAATTATTG
GGNGTAAAGAGTTTGTAGGCGGNTTGTNAAGTNGGTTGTGAAAGCCCGGGGNTTAACCCCGGGTTTGCAGTTGAT
ACGGGCAGGNTAGAGTTCGGTAGGGGAGATNGGAATTCCTGGTGTAGCGGTGAAATGCGCAGATATCAGGAGGAA
CACCGGTGGCGAAGGCGGATCTCTGGGCCATTACTGACGCTGAGGAGCGAAAGCGTGGGGAGCGAACAGGATTAG
ATACCCTGGTAGTCCACGCCGTAAACGGTGGGAAC TAGGTGTTGGCGACATTCCACGTCGTGCGTGCCGCAGCTA
ACGCATTAAGTTCCCCGCCTGGGGAGTACGGCCGCAAGGCTAAACTCAAAGGAATTGACGGGGGCCCGCACAAAG
CAGCGGAGCATGTGGCTTAATTCGACGCAACGCGAAGAACCCTTACCAAGGCTTGACATACACCGGAAAGCATCAG
AGATGGTGCCCCCTTGTGGTGGTGTACAGGTGGTGCATGGCTGTCGTGAGCTCGTGTGAGATGTTGGGTT
AAGTCCCACAACGAGCGCAACCCCTTGTCTGTGTTGCCAGCATGCCCTTCGGGGTGATGGGGACTCACAGGAGAC
CGCCGGGGTCAACTCGGAGGAAGGTGGGGACGACGTCAAGTCATCATGCCCTTATGTCTGGGCTGCACACGTG
CTACAATGGCCGGTACAAAGAGCTGCGATACCGTGAGGTGGAGCGAATCTCAAAAAGCCGGTCTCAGTTCGGATT
GGGGTCTGCAACTCGANCCCATGAANTCGGAGTTGCTAATTAATCGCAAAATCAAGCATTGCTGGCGGTGAATAC
GTTCCC

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**Figure 20**

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Figure 21



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